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5.1.6
Compugen Ltd.
GenCore version (c) 1993 - 2003
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OM protein - protein search, using sw model

November 17, 2003, 18:21:08; Search time 21 Seconds (without alignments) 45.795 Million cell updates/sec Run on:

US-09-462-089-1 63 1 EHWSYGLRPG 10 Perfect score: Sequence: BLOSUM62 Gapop 10.0 , Gapext 0.5 Scoring table:

283308 segs, 96168682 residues Searched:

1100 Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 10

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database :

PIR 76:*
1: pir1:*
2: pir2:*
3: pir3:*
1: pir4:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

	Description	gonadoliberin - pi	•	gonadoliberin I -	gonadoliberin - ch	gonadoliberin II -	gonadoliberin - sp	gonadoliberin II -	gonadotropin-relea	gonadoliberin I -	gonadoliberin - se	neuromedin C - lau	neuromedin C - bov	hain	Ig heavy chain CRD	D-amino-acid oxida	Ig mu chain J regi	fatty-acid synthas	neuropeptide calla	litorin 2-Glu - Au	litorin I - Austra	tachykinin I - mig	tachykinin III - m	Ig heavy chain CRD	hypotrehalosemic h	Ig H chain V-D-J r	translation elonga	b	Ig H chain V-D-J r	T-cell receptor ga
SUMMARIES	ID	RHPGG	RHSHG	RHAQ1	A21114	RHAQ2	A61126	B46030	A49187	A46030	RHLMGS	PQ0177	A60647	PT0299	PT0281	S39437	F33932	A60139	D47393	807205	S07204	ECLOIM	ECLQ3M	PT0268	B33995	PH1618	D58503	PT0238	PH1591	G41946
	Length DB	10 1	10 1	10 1	10 2	10	10 1	10 2	10 2	10 2	10 1	•	10 2	9	•	9	10 2	7 2	•	9	9	10 1	10 1	9			9			9
عن	Query Match Ler	95.2	95.2	88.9	74.6	66.7	66.7	66.7		66.7	41.3	39.7	39.7	34.9		33.3	31.7	0	30.2	0	30.2	30.2	30.2	28.6	œ.	٠	7.	27.0	7.	27.0
	Score	09	9	26	47	42	42	42	42	42	26	25	25	22	21	21	20	19	19	19	19	19	19	18	18	17	17	17	17	17
	Result No.	7	7	e	4	S	9	7	6 0	σ	10	11	12	13	14	15	16	17	18	19	20	21	22	23		25	56		28	29

1 EHWSYGLRPG 10 :|||||||||| QHWSYGLRPG 10

ò g RESULT 2

Grandoliberin - sheep C;Species: Ovis orientalis aries, Ovis ammon aries (domestic sheep) C;Dete: 31-Dec-1991 #sequence_revision 31-Dec-1991 #text_change 18-Mar-1997 C;Accession: A93780; A01411 R;Burgus, R.; Butcher, M.; Amoss, M.; Ling, N.; Monahan, M.; Rivier, J.; Fellows, R.; Bl Proc. Natl. Acad. Sci. U.S.A. 69, 278-282, 1972 A;Title: Primary structure of the ovine hypothalamic luteinizing hormone-releasing facts. A;Reference number: A93780; MUD:72094314; PMID:4550508 A;Accession: A93780

sialokinin I - yel	sialokinin II - ye	scyliorhinin I - s	ranatachykinin B -	cytochrome-c oxida	Ig H chain V-D-J r	hypothetical pepti	litorin - Rohde's	cytochrome-c oxida	sperm-activating p	Ig heavy chain DJ	T-cell receptor al	T-cell receptor ga	T-cell receptor be	244K exoantigen -	T-cell receptor be
A49581	B49581	A24867	B61033	877990	PH1633	155382	S07241	S65388	B60589	PH1344	S23370	F49033	PT0726	A33098	PT0526
7	~	7	7	7	7	4	0	7	~	~	~	~	~	~	~
10	10	70	10	10	10	7	σ	10	10	10	10	10	9	7	7
27.0	27.0	27.0	27.0	27.0	27.0	25.4	25.4	25.4	25.4	25.4	25.4	25.4	23.8	23.8	23.8
17 27.0	17 27.0	17 27.0	17 27.0	17 27.0	17 27.0	16 25.4	16 25.4	16 25.4	16 25.4	16 25.4		16 25.4	15 23.8	15 23.8	15 23.8

ALIGNMENTS

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C;Accession: A01411
R;Baba, Y.; Matsuo, H.; Schally, A.V.
Biochem. Biophys. Res. Commun. 44, 459-463, 1971
A;Title: Structure of the porcine LH- and FSH-releasing hormone. II. Confirmation of the A;Reference number: A90172; MUID:72114303; PMID:4946067
                                                                                                                                                                                                                                                                                                                                               A; Molecule type: protein
A; Residues: 1-10 cBAB>
A; Residues: 1-10 cBAB>
A; Residues: 1-10 cBAB>
B; Matsuo, H.; Arimura, A.; Nair, R.M.G.; Schally, A.V.
Biochem. Biophys. Res. Commun. 45, 822-827, 1971
A; Title: Synthesis of the porcine LH- and FSH-releasing hormone by the solid-phase metho A; Reference number: A90176; MIDD:72065376; PMID:4942726
A; Rotents: annotation; synthesis
A; Note: the synthetic and natural hormones have the same physicochemical and biological A; Note: the synthetic and natural hormones have the same physicochemical and biological A; Arimura, A.; Schally, A.V.
Biochem. Biophys Res. Commun. 45, 483-487, 1971
A; Title: On the tryptophan residue in porcine LH and FSH-releasing hormone.
A; Reference number: A90175; MUID:72117544; PMID:4946275
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A; Contents: annotation
A;Note: Trp-3 appears to be essential for biological activity
C;Comment: This hypothalamic hormone stimulates the secretion of both luteinizing and fc
C;Superfamily: gonadoliberin
C;Keywords: amidated carboxyl end; hypothalamus; pyroglutamic acid
F;1/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental
F;10/Modified site: amidated carboxyl end (Gly) #status experimental
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gonadoliberin - pig
C;Species: Sus scrofa domestica (domestic pig)
C;Date: 13-Jul-1981 #sequence_revision 13-Jul-1981 #text_change 18-Mar-1997
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Best Local Similarity 90.0%
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C,Species: Alligator mississippiensis (American alligator)
C,Date: 31-Mar-1993 #sequence_revision 31-Mar-1993 #text_change 18-Mar-1997
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Best Local Similarity 60.vv,
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1 QHWSHGWYPG 10
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QHWSHGWYPG 10
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R; Lovejoy, D.A.; Fischer, W.H.; Parker, D.B.; MCROry, J.E.; Park, M.; Lance, V.; Swanson R; Lovejoy, D.A.; Fischer, W.H.; Parker, D.B.; MCROry, J.E.; Park, M.; Lance, V.; Swanson Regul. Pept. 3, 105-116, 1991
A; Title: Primary structure of two forms of gonadotropin-releasing hormone from brains of A; Reference number: A60066; MUD:91352338; PMID:1882082
A; Accession: A60066
A; Molocule type: Portein
A; Residues: 1-10 < LOV-
C; Superfemily: gonadollberin
C; Superfemily: gonadollberin
C; Keywords: amidated carboxyl end; hormone; hypothalamus; pyroglutamic acid
F; 1/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental
F; 10/Modified site: amidated carboxyl end (Gly) #status experimental
                        A; Note: the natural and synthetic hormones have the same biological activity C; Comment: This hypothalamic hormone stimulates the secretion of both luteinizing and C; Superfamily: gonadoliberin (C; Superfamily: gonadoliberin (C; Keywords: amidated carboxyl end; hormone; hypothalamus; pyroglutamic acid F; I, Modified site: pyrrolidone carboxylic acid (Gln) #status experimental F; 10/Modified site: amidated carboxyl end (Gly) #status experimental
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               gonadoliberin I - American alligator
N;Alternate names: gonadotropin-releasing hormone I
C;Species: Alligator mississippiensis (American alligator)
C;Date: 31-Mar-1993 #sequence_revision 31-Mar-1993 #text_change 18-Mar-1997
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A;Title: Characterization of a teleost gonadotropin-releasing hormone. A;Reference number: A21114; WUID:83195140; PMID:6341999
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                                                                                                                                                                                                           Score 60; DB 1; Length 10;
Pred. No. 0.00028;
1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       88.9%; Score 56; DB 1; Length 10; 80.0%; Pred. No. 0.0013; ive 2; Mismatches 0; Indels
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N;Alternate names: gonadotropin-releasing hormone
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C;Species: Oncorhynchus keta (chum salmon)
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Best Local Similarity 80.0
Matches 8; Conservative
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A; Status: preliminary
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C;Accession: B60066
R;Lovejoy, D.A.; Fischer, W.H.; Parker, D.B.; McRory, J.E.; Park, M.; Lance, V.; Swanson sequl. Pept. 33, 105-116, 1991
A;Title: Primary structure of two forms of gonadotropin releasing hormone from brains of A;Reference number: A60066; MUID:91352338; PMID:1882082
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         C,Accession: A61126
R;Lovejoy, D.A.; Sherwood, N.M.; Fischer, W.H.; Jackson, B.C.; Rivier, J.B.; Lee, T.
R;Lovejoy, D.A.; Sherwood, N.M.; Fischer, W.H.; Jackson, B.C.; Rivier, J.B.; Lee, T.
A;Title: Primary structure of gonadotropin-releasing hormone from the brain of a holocep A;Reference number: A61126; MUID:91340067; PMID:1678723
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Aritle: Distinct sequence of gonadotropin-releasing hormone (GnRH) in dogfish brain proc. A;Reference number: A46030; MUID:92335300; PMID:1631133
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  gonadoliberin - spotted ratfish
NyAlernate names: gonadotropin-releasing hormone
C.Species: Hydrolagus colliei (spotted raffish)
C.Species: 26-May-1994 #sequence_revision 26-May-1994 #text_change 18-Mar-1997
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A,Molecule type: protein
A,Rolacule type: Drotein
C,Superfamily: gonadoliberin
C,Keywords: hormone; pyroglutamic acid
E,1/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental
                                                                                                                                                                                                                       A;Accession: B60066
A;Aolacule type: protein
A;Residues: 1-10 <100v.
C;Superfamily: gonadoliberin
C;Stwperfamily: gonadoliberin
C;Keywords: amidated carboxyl end; hormone; hypothalamus; pyroglutamic ac.
F;1/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental
F;10/Modified site: amidated carboxyl end (Gly) #status experimental
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Pred. No. 0.32;
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C;Accession: PQ0177
R;Conlon, J.M.; O'Harte, F.; Vaudry, H.
Rabcchem. Biophys. Res. Commun. 178, 525-530, 1991
A;Title: Primary structures of the bombesin-like neuropeptides in frog brain show that b. A;Reference number: PQ0177; WUID:91315477; PMID:1859413
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    C;Accession: Å60647

R;Lemaire, S.; Trifaro, J.M.; Chouinard, L.; Cecyre, D.; Dessureault, J.; Mercier, P.; D

Reptides 10, 355-360, 1989

A;Title: Structural identification, subcellular localization and secretion of bovine adr

A;Reference number: A60647; MUID:89331342; PMID:2755876
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A; Note: this neuropeptide was purified from secretory granules of cells in the adrenal m
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         neuromedin C - laughing frog
C;Species: Rana ridibunda (laughing frog)
C;Date: 23-Nov-1991 #sequence_revision 23-Nov-1991 #text_change 11-Jan-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          C;Species: Bos primigenius taurus (cattle)
C;Date: 14-May-1993 #sequence_revision 14-May-1993 #text_change 07-May-1999
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                                                                                                                                                                                                                                                Gaps
                             A, Residues: 1-10 <SHE>
C;Comment: This hormone was isolated from the brain.
C;Comment: This hormone was isolated from the brain.
C;Superfamily: gonadoliberin
C;Keywords: amidated carboxyl end; hormone; pyroglutamic acid
F;1/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental
F;10/Modified site: amidated carboxyl end (Gly) #status experimental
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C.Superfamily: gastrin-releasing peptide
C.Keywords: amidated carboxyl end
F;10/Modified site: amidated carboxyl end (Met) #status predicted
                                                                                                                                                                        Score 26; DB 1; Length 10, Pred. No. 1.78+02;
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Pred. No. 2.4e+02;
1; Mismatches 1;
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60.0%; Pred. No. 2.4e+02;
tive 1; Mismatches 1;
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                                                                                                                                                                                              41.3%;
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Best Local Similarity 60.v.
.h.e 3; Conservative
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Best Local Similarity 40.0
'... 4; Conservative
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         A; Molecule type: protein
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HWAVG 7
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R;Lovejoy, D.A.; Fischer, W.H.; Ngamvongchon, S.; Craig, A.G.; Nahorniak, C.S.; Peter, R
Proc. Natl. Acad. Sci. U.S.A. 89, 6373-6377, 1992
A;Title: Distinct sequence of gonadotropin-releasing hormone (GnRH) in dogfish brain pro
A;Reference number: A46030; MUID:92335300; PMID:1631133
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R;Sower, S.A.; Chiang, Y.C.; Lovas, S.; Conlon, J.M.
Bidocrinology 132, 1125-1131, 1993
A;Tille: Primary structure and biological activity of a third gonadotropin-releasing A;Reference number: A49187; MUID:93178316; PMID:8440174
A;Accession: A49187
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N;Alternate names: gonadotropin releasing hormone (GnRH)
N;Alternate names: gonadotropin releasing hormone (GnRH)
C;Species: Petromyzon marinus (sea lamprey)
C;Date: 17-Mar-1987 #sequence_revision 17-Mar-1987 #text_change 18-Mar-1997
C;Accession: A01412
R;Sherwood, N.M.; Sower, S.A.; Marshak, D.R.; Fraser, B.A.; Brownstein, M.J.
Biol. Chem. 251, 4812-4819, 1986
A;Title: Primary structure of gonadotropin-releasing hormone from lamprey brain.
A;Reference number: A01412; MUID:86168192; PMID:3514603
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Date: 06-Jan-1995 #sequence_revision 06-Jan-1995 #text_change 16-Dec-1998
                                                                                                                                                                                                                                                          Species: Petromyzon marinus (sea lamprey)
State: 19-Dec-1993 #sequence_revision 18-Nov-1994 #text_change 03-Mar-1995
Accession: A49187
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A;Residues: 1-10 <LOV>
C;Keywords: hormone; pyroglutamic acid
F;1/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental
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A;Residues: 1-10 <SOW>
A;Experimental source: brain
A;Note: sequence extracted from NCBI backbone (NCBIP:126381)
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60.0%; Pred. No. 0.32;
cive 2; Mismatches
         Pred. No. 0.32;
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Pred. No. 0.32;
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                                2; Mismatches
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       60.08;
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N'Alternate names
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Best Local Similarity 60.0
Matches 6; Conservative
                                6; Conservative
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QHWSHGWYPG 10
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C;Species: Trigonopsis variabilis
C;Species: Trigonopsis variabilis
C;Date: 19-Mar-1997 #sequence_revision 05-Dec-1997 #text_change 07-May-1999
C;Accession: 339437
R;Schraeder, T.; Andreesen, J.R.
Ebur. J. Biochem. 218, 735-744, 1993
A;Title: Evidence for the functional importance of Cys298 in D-amino acid oxidase from TA;Reference number: S39437
A;Reference number: S39437
A;Reference protein
A;Residues: 1-9 <SCH-
A;Residues: Protein
C;Species: Homo sapiens (man)
C;Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change 16-Aug-1996
C;Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change 16-Aug-1996
C;Accession: PT029
R;Yamada, M; Wasserman, R.; Reichard, B.A.; Shane, S.; Caton, A.J.; Rovera, G.
J. Exp. Med. 173, 395-407, 1991
J. Exp. Med. 173, 395-407, 1991
A;Reference number: PT0222; MUID:91108337; PMID:1899102
A;Accession: PT0299
A;Molecule type: DNA
A;Molecule type: DNA
A;Residues: 1-9 <XAM>A;Residues: 1-9 <XAM>A;Residue
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C;Species: Homo sapiens (man)
C;Species: Homo sapiens (man)
C;Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change 16-Aug-1996
C;Accession: PT0281
R;Yamada, M: Wasserman, R.; Reichard, B.A.; Shane, S.; Caton, A.J.; Rovera, G.
J. Exp. Med. 173, 395-407, 1991
A;Title: Preferential utilization of specific immunoglobulin heavy chain diversity and A;Accession: PT0281
A;Accession: PT0281
A;Molecule type: DNA
A;Molecule type: DNA
A;Molecule type: DNA
A;Residues: 1-5 < YAM>
A;Residues: 1-5 < YAM>
A;Residues: 1-5 < YAM>
A;Residues: a lymphocyte
C;Keywords: heterotetramer; immunoglobulin
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16-OCT-2001 (Rel. 40, Created)
16-OCT-2001 (Rel. 40, Last sequence update)
28-FEB-2003 (Rel. 41, Last annotation update)
Gonadoliberin I (Gonadotropin-releasing hormone I) (GnRH-I) (LH-RH)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Clupea pallasii (Pacific herring).
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Actinopterygii, Neopterygii, Teleostei, Clupeomorpha, Clupeidae,
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SPECIES—C. Pallasii; PubMed=10650929;
Carolafeld J., Powell J.F., Park M., Fischer W.H., Craig A.G.,
Carolafeld J., Powell J.F., Park M., Fischer W.H., Craig A.G.,
Chang J.P., Rivier J.E., Sherwood N.M.;
"Primary structure and function of three gonadotropin-releasing
hormones, including a novel form, from an ancient teleost, herring.";
Endocrinology 141:505-512(2000).
-- FUNCTION: Stimulates the secretion of gonadotropins; it stimulates
the secretion of both luteinizing and follicle-stimulating
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P20367; P81751;
01-FEB-1991 (Rel. 17, Created)
01-FEB-1993 (Rel. 17, Last sequence update)
28-FEB-2003 (Rel. 41, Last annotation update)
Gonadoliberin III (Gonadotropin-releasing hormone III) (GRRH-III) (LH-
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Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;

Actinopterygii; Neopterygii; Teleostei; Buteleostei;

Protacanthopterygii; Salmoniformes; Salmonidae; Oncorhynchus.

NCBI_TaxID=8018, 30724;
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MEDLINE=81195140; PubMed=6341999;
MEDLINE=81195140; L., Brownstein M., Spiess J., Rivier J., Vale 'Sherwood N., Eiden L., Brownstein M., Spiess J., Rivier J., Vale 'Characterization of a teleost gonadotropin-releasing hormone."; proc. Natl. Acad. Sci. U.S.A. 80:2794-2798(1983).
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2; Mismatches
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!- SUBCELLULAR LOCATION: Secreted
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Best Local Similarity 70.0%;
Matches 7; Conservative
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InterPro; IPR002012; GnRH.
Pfam; PF00446; GnRH; 1.
PROSITE; PS00473; GNRH; 1.
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                                            NCBI TaxID=30724;
                                                                                                                        SEQUENCE,
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D GON2_CHICK STANDARD; PRT; 10 AA.

D GON2_CHICK STANDARD; PRT; 10 AA.

C P37043; P270408; P81750;

T 01-FEB-1991 (Rel. 17, Created)

T 28-FEB-2003 (Rel. 41, Last annotation update)

G Gonadoliberin II (Gonadortropin-releasing hormone II) (GnRH-II)

E Gonadoliberin II) (American alligator),

Alligator missisabpiplensis (American alligator),

Alligator missisabpiplensis (American alligator),

S Qualus acanthias (Spiny dogfish),

Hydrolagus colliei (Spotted ratfish) (Pacific ratfish), and

C Hydrolagus (Spiny dogfish),

C Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

C Archosauria; Aves; Neognathae; Galliformes; Phasianidae; Phasianinae;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        "Identification of the second gonadotropin-releasing hormone in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SPECIES-S. acanthias; TISSUE=Brain; MEDLINE=92335300; PubMed=1631133; Lovejoy D.A., Fischer W.H., Ngamvongchon S., Craig A.G., Nahorniak C.S., Peter R.E., Rivier J.E., Sherwood N.M.; "Distinct sequence of gonadotropin-releasing hormone (GnRH) in dogfish brain provides insight into GnRH evolution."; Proc. Natl. Acad. Sci. U.S.A. 89:6373-6377(1992).
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PYRROLIDONE CARBOXYLIC ACID
                                                                                                                                                                                                         2; Indels
                                                                                                                                 Length
                                                                284B3233786B45A3 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            [5]
SEQUENCE, AND FUNCTION.
SPECIES=C.pallasi; TISSUE-Brain, and Pituitary;
MEDLINE-20114351; Pubmed=10650929;
                                                                                                                                     Score 47; DB 1;
Pred. No. 0.014;
.; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SPECIES=A.mississippiensis; TISSUE=Brain;
MEDLINE=91352338; PubMed=1882082;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NCBI_TaxID=9031, 8496, 7797, 7873, 30724;
                                   AMIDATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SPECIES=Chicken; TISSUE=Hypothalamus;
MEDLINE=84222059; PubMed=6427779;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDLINE=91340067; PubMed=1678723;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE.
SPECIES=H.colliei; TISSUE=Brain;
1
10
1230 MW;
                                                                                                                                     74.6%;
70.0%;
                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                             EHWSYGLRPG 10
                                                                                                                                                                                                                                                                                                                                                  OHWSYGWLPG 10
1
10
10 AA;
                                                                                                                                                              Local Similarity
es 7; Conserv
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10 AA

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STANDARD;
                                                                                                                                                                                                                                                                                                                                                  Best Local Similarity
                                                                                                       NCBI_TaxID=7757;
                                                             (Luliberin III)
                                                                                                                                      TISSUE=Brain;
                                                                                                                                                                                                                        hormones.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (Luliberin I)
GON3_PETMA
P30948;
01-JUL-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                           CHEPR
                                                                                                                                                                                                                                                                                                          MOD_RES
SEQUENCE
                                                                                                                                                                                                                                                                                                                                         Query Match
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                                                                                                                            SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                              GON1_CHEPR
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0
      Gaps
                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                    Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Chondrichthyes;
Elasmobranchii; Squalea; Squaloidei; Squalidae; Squalus.
 Carolsfeld J., Powell J.F.F., Park M., Fischer W.H., Craig A.G.,
                                                                                                                                                                                                                       ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDLINES-9233300; PubMed=1631133; MEDLINES-9233300; PubMed=1631133; Media 9. Craig A.G., Lovejoy D.A., Fischer W.H., Ngamvongchon S., Craig A.G., Nahorniak C.S., Peter R.E., Rivier J.E., Sherwood N.M.; Distinct sequence of gonadorropin-releasing hormone (GnRH) in dogitab brain provides insight into GRHR evolution."; Proc. Natl. Acad. Sci. U.S.A. 89:6373-6377 (1992).

-: FUNCTION: STIMULATES THE SECRETION OF GONADOTROPINS.
-: SUBCELLULAR LOCATION: Secreted.
-: SIMILARITY: Belongs to the GNRH family.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ö
                                                                                                                                             Hormone; Amidation; Hypothalamus; Pyrrolidone carboxylic acid.

NOD RES 1 1 PRRROLIDONE CARBOXYLIC ACID.

AMIDATION AMIDATION 2860UENCE 10 AA; 1254 MW; 284B2E437871F5A3 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Pfam; PF00446; GnRH; 1.
PROSITE; PS00473; GNRH; 1.
Hormone; Amidation; Hypothalamus; Pyrrolidone carboxylic acid.
MOD_RES 1 1 PYRROLIDONE CARBOXYLIC ACID.
MOD_RES 10 AMIDATION.
SEQÜENCE 10 AA; 1204 MW; 284832337871F5A3 CRC64;
                                                                                                                                                                                                                                                                                                                                        01-AUG-1992 (Rel. 23, Created)
01-AUG-1992 (Rel. 23, Last sequence update)
28-FEB-2003 (Rel. 41, Last annotation update)
Gonadoliberin (Gonadotropin-releasing hormone) (GRRH) (LH-RH)
                                                                                                                                                                                                   DB 1; Length 10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 1; Length 10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2; Indels
                                                                                                                                                                                                                       2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 42; DB 1;
Pred. No. 0.1;
2; Mismatches
                                                                                                                                                                                                                                                                                                                    10 AA
                                                                                                                                                                                                           Pred. No. 0.1;
2; Mismatches
                                                                                                                                                                                                  Score 42;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                             Squalus acanthias (Spiny dogfish)
                                                                                                                                                                                                  66.78;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             66.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         InterPro; IPR002012; GnRH.
                                                                                                                                                                                                                       6; Conservative
                                                                                                                                                                                                                                                                                                                    STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 EHWSYGLRPG 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 OHWSHGWLPG 10
                                                                                                                                                                                                                                                        1 EHWSYGLRPG 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
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Best Local Similarity
Matches 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                           NCBI_TaxID=7797;
                                                                                                                                                                                                                                                                                                                                                                                   Luliberin).
                                                                                                                                                                                                                                                                                                                    SOUAC
                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE
                                                                                                                                                                                                                                                                                                                              P27429
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ID GONL 9
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GON3_PETMA

RESULT 6

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-!- SUBCELLULAR LOCATION: Secreted.
-!- TISSUE SPECIFICITY: GNRH NEURONS LIE WITHIN BLOOD SINUSES CLOSE TO TISSUE SPECIFICITY: GNRH NEURONS LIE WITHIN BLOOD SINUSES CLOSE TO THE GONODUCTS AND GONADS IN BOTH JUVENILES AND ADULTS, IMPLYING THAT THE NEUROPEPTIDE IS RELEASED INTO THE BLOODSTREAM.
-!- MASS SPECTROMETRY: MW=1246.56; METHOD=MALDI.
-!- SINILARITY: Belongs to the GNRH family.
InterPro; IPR002012; GNRH.
PF00446; GNRH; 1.
                                                                                                                                                                                                                                               MEDLINE=93178316; PubMed=8440174;

Sower S.A., Chiang Y.-C., Lovas S., Conlon J.M.;

Sower S.A., Chiang Y.-C., Lovas S., Conlon J.M.;

"Primary structure and biological activity of a third gonadotropin-

releasing hormone from lamprey brain.";

Endocrinology 132:1125-1131(1993).

-! FUNCTION: Stimulates the secretion of gonadotropins; it stimulates

the secretion of both lutelinizing and follicle-stimulating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Powell J.F.E., Reska-Skinner S.M., Prakash M.O., Fischer W.H.,
Park M., Rivier J.E., Craig A.G., Mackie G.O., Sherwood N.M.;
"Two new forms of gonadotropin-releasing hormone in a protochordate
and the evolutionary implications.";
Proc. Natl. Acad. Sci. U.S.A. 93:10461-10464 (1996).

- FUNCTION: Stimulates the secretion of gonadotropins; it stimulates
the secretion of both luteinizing and follicle-stimulating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Chelyosoma productum.
Bukaryota; Metazoa; Chordata; Urochordata; Ascidiacea; Enterogona;
Phlebobranchia; Corellidae; Chelyosoma.
MCBI_TaxID=71177;
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Hyperoartia,
Petromyzontiformes, Petromyzontidae, Petromyzon.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ..
0
                                                                                                                                                                                                                                                                                                                                                                                                    01-NOV-1997 (Rel. 35, Created)
01-NOV-1997 (Rel. 35, Last sequence update)
28-FEB-2003 (Rel. 41, Last annotation update)
Gonadoliberin I (Gonadotropin-releasing hormone I) (GnRH-I)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 39; DB 1; Length 10; Pred. No. 0.36;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AMIDATION.
284B36237AA1F5A3 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              10 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       3; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDLINE=96413669; PubMed=8816823;
                                                                                                     Petromyzon marinus (Sea lamprey).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                61.9%;
50.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     10 AA; 1277 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         5; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 EHWSYGLRPG 10
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HWAVG
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P23260;
                                                                                                                                                                                                                       MOD_RES
SEQUENCE
                                       SEQUENCE
                                                                                                                                                                                                                                                     Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                  -:- SUBCELLULAR LOCATION: Secreted.
-:- TISSUE SPECIFICITY: GNRH NEURONS LIE WITHIN BLOOD SINUSES CLOSE TO THE GONDUCTS AND CONADS IN BOTH UVENILES AND ADULTS, IMPLYING THAT THE NEUROPEPTIDE IS RELEASED INTO THE BLOODSTREAM.
-:- SMILLARITY: Belongs to the GNRH Family.
InterPro; IPR002012; GnRH.
PROSITE; PS00473; GNRH; 1.
                                                                                                                                                                                                                                                                                                             Powell J.F.F., Reska-Skinner S.M., Prakash M.O., Fischer W.H.,
Park M., Rivier J.E., Craig A.G., Mackie G.O., Sherwood N.M.;
"Two new forms of gonadotropin-releasing hormone in a protochordate
and the evolutionary implications.";
Proc. Natl. Acad. Sci. U.S.A. 93:10461-10464 (1996).

-!- FUNCTION: Stimulates the secretion of gonadotropins; it stimulates
the secretion of both luteinizing and follicle-stimulating
                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                  Bukaryota, Metazoa, Chordata, Urochordata, Ascidiacea, Enterogona,
Phlebobranchia, Corellidae, Chelyosoma.
NCBI_TaxID=71177;
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                                                                                                                                                                                         01-NOV-1997 (Rel. 35, Created)
01-NOV-1997 (Rel. 35, Last sequence update)
28-FEB-2003 (Rel. 41, Last annotation update)
Gonadoliberin II (Gonadotropin-releasing hormone II) (GRH-II)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             20-MAR-1987 (Rel. 04, Created)
20-MAR-1987 (Rel. 04, Last sequence update)
28-F8B-2003 (Rel. 41, Last annotation update)
Gonadoliberin I (Gonadotropin-releasing hormone I) (GnRH-I)
        Score 30; DB 1; Length 10;
Pred. No. 14;
1; Mismatches 4; Indels
                                                          Length 10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PYRROLIDONE CARBOXYLIC ACID.
                                                                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AMIDATION (BY SIMILARITY).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            284B38D1EEB735A3 CRC64;
                                                         Score 34; DB 1;
Pred. No. 2.7;
2; Mismatches 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           10 AA.
                                                                                                                                                                       10 AA
                                                                                                                                                                                                                                                                                                                                                                                             -!- SUBUNIT: Homodimer; disulfide-linked.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        INTERCHAIN.
                                                                                                                                                                     PRT;
                                                                                                                                                                                                                                                                                                    MEDLINE=96413669; PubMed=8816823;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (Laliberin I).
Petromyzon marinus (Sea lamprey)
                                                          54.0%;
50.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                47.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            10 AA; 1135 MW;
 PS00473; GNRH; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity 50.0
                                                                              5, Conservative
                                                                                                                                                                      STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STANDARD;
                                                                                                   1 EHWSYGLRPG 10
                                                                                                                     1 QHWSDYFKPG 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 EHWSYGLRPG 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 QHWSLCHAPG 10
                                                                                                                                                                                                                                           Chelyosoma productum
                                                                     Best Local Similarity
                                                                                                                                                                                                                                 (Luliberin II)
                                                                                                                                                                                                                                                                                                                                                                                     hormones.
                                                                                                                                                                      CHEPR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PETMA
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SEQUENCE
          Hormone;
MOD_RES
MOD_RES
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 PROSITE;
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P04378;
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                                                                              Matches
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Sherwood N.M., Sower S.A., Marshak D.R., Fraser B.A., Brownstein M.J., "Primary structure of gonadotropin-releasing hormone from lamprey
                                                                                                                                                                                                                              gonadotropins; it stimulates follicle-stimulating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Rana ridibunda (Laughing frog) (Marsh frog).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Amphibia; Batrachia; Anura; Neobatrachia; Ranoidea; Ranidae; Rana.
VCBI_TaxID=8406;
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Hyperoartia, Petromyzontiformes, Petromyzontidae, Petromyzon.
NCBI_TaxID=7757;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ..
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0
                                                                                                                                                                                                                                                                                                                                                                                                                Hormone, Amidation, Hypothalamus, Pyrrolidone carboxylic acid. MOD_RES 1 1 PYRROLIDONE CARBOXYLIC ACID.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 25; DB 1; Length 10;
Pred. No. 1.1e+02; .
1; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1E4B36237B1735AB CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AMIDATION.
F81FBAE862CDC371 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 26; DB 1;
Pred. No. 71;
3; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     01-NOV-1991 (Rel. 20, Created)
01-NOV-1991 (Rel. 20, Last sequence update)
28-FEB-2003 (Rel. 41, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               10 AA
                                                                                                                                                                                                                                                                                          -!- SUBCELLULAR LOCATION: Secreted.
-!- SIMILARITY: Belongs to the GNRH family.
PIR; A01412; RHIMGS.
InterPro. IPR002012; GNRH.
                                                                                                                                                                                                       J. Biol. Chem. 261:4812-4819(1986).
                                                                                                       TISSUE=Brain;
MEDLINE=86168192; PubMed=3514603;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PIR; PQ0177; PQ0177.
InterPro; IPR000874; Bombesin.
Pfam; PF02044; Bombesin; I.
PROSITE; PS00257; BOMBESIN; I.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       n
Similarity 60.0%;
3; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE 10 AA; 1094 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    41.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         40.08;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Bombesin family; Amidation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            10 AA; 1244 MW;
                                                                                                                                                                                                                                                                                                                                                                Pfam; PF00446; GnRH; 1.
PROSITE; PS00473; GNRH; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 EHWSYGLRPG 10
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Best Local Similarity
Matches 3; Conserv
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"ISOJATION and identification of multiple neuropeptides of the alactostatin superfamily in the shore crab Carcinus maenas."; Eur. J. Biochem. 250:727-734(1997).
-!- FUNCTION: MAY ACT AS A NEUROTRANISMITTER OR NEUROMODULATOR.
-!- SIMILARITY: BELONGS TO THE ALLATOSTATIN FAMILY.
Neuropeptide, Amidation, Multigene family.
MOD RES S AMIDATION (POTENTIAL).
SEQUENCE 5 AA, 586 MW; 672879D5AB300000 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         INDIAR AND AND IDENTIFICATION OF MULTIPLE NEUROPEPTIDES OF THE ALLACOSTATION AND ADDRESS."; EUR. J. BLOCHEM. 250:77-734(1997).

- FUNCTION: MAX ACT AS A NEUROTRANSMITTER OR NEUROMODULATOR.

- SIMILARITY: BELONGS TO THE ALLATOSTATIN PAMILY.
                                                                                                                                                                                                                            TISSUE=Cerebral ganglion, and Thoracic ganglion;
MEDLINE=98121193; PubMed=9461295;
Duve H., Johnsen A.H., Maestro J.-L., Scott A.G., Jaros P.P.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Duve H., Johnsen A.H., Maestro J.-L., Scott A.G., Jaros P.P.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Bukaryota; Metazoa; Arthropoda; Crustacea; Malacostraca;
Bumalacostraca; Bucarida; Decapoda; Pleocyemata; Brachyura;
Bubrachyura; Portunoidea; Portunidae; Carcinus.
                                                                                                             Carcinus maenas (Common shore crab) (Green crab).
Bukaryota; Metazoa; Arthropoda; Crustacea; Malacostraca;
Bumalacostraca; Eucarida; Decapoda; Pleocyemata; Brachyura;
Bubrachyura; Portunoidea; Portunidae; Carcinus.
                                                                                                                                                                                                                                                                                                                                                                                                                                              30.2%; Score 19; DB 1; Length 5; 60.0%; Pred. No. 1.3e+05; trive 2; Mismatches 0; Indels
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MOD RES 7 7 7 AMIDATION (POTENTIAL).

SEQUENCE 7 AA; 770 MW; 672879CDCB5DDB70 CRC64;
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Carcinus maenas (Common shore crab) (Green crab)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                30.2%; Score 19; DB 1; I 42.9%; Pred. No. 1.3e+05; trive 3; Mismatches 1;
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(Rel. 39, Last sequence update)
(Rel. 39, Last annotation update)
                                              (Rel. 39, Created)
(Rel. 39, Last sequence update)
(Rel. 39, Last annotation update)
                 5 AA.
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1 YSFGL 5
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MEDLINE=98121193; PubMed=9461295;
Duve H., Johnsen A.H., Maestro J.-L., Scott A.G., Jaros P.P.,
                                                                                                                                            Eukaryota, Metazoa, Arthropoda, Hexapoda, Insecta, Pterygota, Neoptera, Endopterygota, Lepidoptera, Glossata, Ditrysia, Tortricoidea, Tortricidae, Olethreutinae, Cydia.
                                                                                                                                                                                                                                                                            Duve H., Johnsen A.H., Maestro J.-L., Scott A.G., Winstanley Davey M., East P.D., Thorpe A.;
"Lepidopteran peptides of the allatostatin superfamily.";
Peptides 18:1301-1309(1997).
-!- SIMILARITY: BELONGS TO THE ALLATOSTATIN FAMILY.
Neuropeptide; Amidation.
MOD RES 8

                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Carcinus maenas (Common shore crab) (Green crab).
Eukaryota; Metazoa; Arthropoda; Crustacea; Malacostraca;
Eumalacostraca; Eucarida; Decapoda; Pleocyemata; Brachyura;
Eubrachyura; Portunoidea; Portunidae; Carcinus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Indels
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        MOD RES
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        8
        AMIDATION

        SEQUENCE
        8 AA; 934 MW; C82879C45B51F775 CRC64;

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30-MAY-2000 (Rel. 39, Last sequence update)
30-MAY-2000 (Rel. 39, Last annotation update)
                                          P82152;
30-MAY-2000 (Rel. 39, Created)
30-MAY-2000 (Rel. 39, Last sequence update)
30-MAY-2000 (Rel. 39, Last annotation update)
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                                                                                                               Cydiastatin 1.
Cydia pomonella (Codling moth).
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Best Local Similarity 50.0
Matches 3; Conservative
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                              STANDARD;
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Matches 4; Conserv
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3 HYNFGL 8
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YSYGL 8
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TISSUE=Larva;
                              CYDPO
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Length 7; 1; Indels

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RESULT 15
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DT 30-MAY-2000 (Rel. 39, Last sequence update)

DT 30-MAY-2000 (Rel. 39, Last annotation update)

DT 30-MAY-2000 (Rel. 39, Last annotation update)

DG Carcinustatin 3.

Carcinus maenas (Common shore crab) (Green crab)

CE Enwalacostraca; Eucarida; Decapoda; Pleocyemata;

DC Enwalacostraca; Eucarida; Decapoda; Pleocyemata;

CE Enwalacostraca; Eucarida; Decapoda; Pleocyemata;

NOTEL TAXID=6759;

RN (TISSUB-Creerbral ganglion, and Thoracic ganglion;

RN MEDLINE=98121193; Pubmed=9461295;

RA Thorpe A.; Johnsen A.H., Maestro J.-L., Scott A.G., Jaros P.P.,

RA Thorpe A.;

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Search completed: November 17, 2003, 18:22:18 Job time : 11 secs

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O9prid Tepisocreus
O9tr47 bos taurus
O9tr47 bos taurus
O94d5 varanus oli
O69347 herpes simp
O85406 coxiella bu
P70243 mus musculu
O9xalo capra hircu
Q8tg88 pleurotus o
                                                  Q19888 homo sapien
094vb2 varanus sal
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094vb5 varanus sal
064971 alfalfa mos
09byf9 homo sapien
09twv0 anthopleura
09mw43 homo sapien
095xv6 varanus pil
06530 mus spretus
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Maccy J.R., Strasburg J.L., Brisson J.A., Vredenburg V.T.,
Jennings M., Larson A.;
Molecular Phylogenetics of Western North American Frogs of the Rana
boylii Species Group.";
Mol. Phylogenet. Evol. 19:131-143(2001).
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Amphibia; Batrachia; Anura; Neobatrachia; Ranoidea; Ranidae; Rana.
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Pred. No. 4.1e+03;
1; Mismatches 1; Indels
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01-DEC-2001 (TrEMBLrel. 19, Last sequence
01-DEC-2001 (TrEMBLrel. 19, Last annotation
Cytochrome c oxidase subunit I (Fragment)
                                                                                                                                                                                                                                                                                                                                                                                                                                           ALIGNMENTS
             090TR48
090VE6
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50.0%;
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10 AA; 1335 MW;
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                                                     NCBI_TaxID=160500;
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SEQUENCE
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094px5 felis silve
094px7 felis silve
094px6 felis libyc
08wbx4 diadema mex
08 fer ficedula al
08 fer ficedu
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(c) 1993 - 2003 Compugen Ltd
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No.
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STRAIN-1, 2, 7, 12, 16, 17, and 110;
Randi E., Pierpaoli M., Beaumont M., Ragni B., Sforzi A.;
Randi E., Pierpaoli M., Beaumont M., Ragni B., Sforzi A.;
"Genetic identification of wild and domestic cats (Felis silvestris),
Genetic identification of wild and domestic cats (Relis silvestris),
Submitted (MAR-2001) to the EMBL/GenBank/DDBJ databases.
EMBL, AJ409129; CAC41030.1; --
EMBL, AJ409130; CAC41030.1; --
EMBL, AJ409131; CAC41036.1; --
EMBL, AJ409131; CAC41036.1; --
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STRAIN=40, 1, 2, and 7;
STRAIN=40, 1, 2, and 7;
Randi E., Pierpaoli M., Beaumont M., Ragni B., Sforzi A.;
"Genetic identification of wild and domestic cats (Felis silvestris) and their hybrida using Bayesian clustering methods.";
Submitted (MAR.2001) to the EMBL/GenBank/DDBJ databases.
EMBL; AJ409135; CAC41068.1; --
EMBL, AJ409140; CAC41069.1; --
EMBL, AJ409142; CAC41069.1; --
EMBL, AJ409144; CAC41075.1; --
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Mammalia; Eutheria; Carnivora; Fissipedia; Felidae; Felis.
NCBI_TaxID=61377;
                                                                                                                                                                             Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Carnivora, Fissipedia, Felidae, Felis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1; Indels
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01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
Cytochrome c oxidase subunit II (Fragment).
                                                       Last sequence update)
Last annotation update)
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75.0%; Pred. No. 8.3e+05;
tive 0; Mismatches 1;
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                                                                                               Cytochrome c oxidase subunit II (Fragment)
                                     Created)
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                                                                                                                                        Felis silvestris catus (Cat).
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01-DEC-2001 (TrEMBLrel. 19,
01-DEC-2001 (TrEMBLrel. 19,
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Best Local Similarity 75.0
Matches 3; Conservative
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PRELIMINARY;
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NON TER
SEQUENCE 8 A
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                  Q94PX7
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EMBL;
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Q94PX6
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                                                                                                                                                                                                                                                                             "Evidence for insufficient chondrocytic differentiation during repair of full-thickness defects of articular cartilage.";
Matrix Biol. 15:39-47(1996).
EMBL; S83371; AAD14433.1; -.
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Randi E., Pierpaoli M., Beaumont M., Ragni B., Sforzi A.;

"Genetic identification of wild and domestic cats (Pelis silvestris),

"Genetic identification of wild and domestic cats (Pelis silvestris),

Submitted (MAR-2001) to the EMBL/GenBank/DDBJ databases.

EMBL, AJ409136; ACA410541; --

EMBL, AJ409139; ACA410541; --

EMBL, AJ409139; ACA41060.1; --

EMBL, AJ409141; CAC41060.1; --

EMBL, AJ409141; CAC41060.1; --

EMBL, AJ409141; CAC41060.1; --

EMBL, AJ409141; CAC41060.1; --
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Mammalia, Eutheria, Lagomorpha, Leporidae, Oryctolagus.
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Mammalia, Eutheria, Carnivora, Fissipedia, Felidae, Felis.
                                                                                                                                                                                                  SEQUENCE FROM N.A.
MEDLINE-96377339; PubMed-8783186;
Metsaranta M., Kujala U.M., Pelliniemi L., Osterman H., Aho H.,
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8.3e+05;
hes 0; Indels
                01-JUL-1997 (TrEMBLrel. 04, Created)
01-JUL-1997 (TrEMBLrel. 04, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
Pro alpha 1 type III collagen protein (Fragment).
Oryctolagus cuniculus (Rabbit).
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SEQŪENCE 8 AA; 1028 MW; B859C7272EA77371 CRC64;
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8 AA; 951 MW; 262685BDC5A3733B CRC64;
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01-DEC-2001 (TrEMBLrel. 19, Last ann
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Best Local Similarity
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Best Local Similarity
Matches 2; Conserv
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                                                                                                                                                             NCBI_TaxID=9986;
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SEQUENCE
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Best Local Similarity
Matches 3; Conserv
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3 YGVLP 7
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ID Q8JJ33
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Archosauria; Aves; Neognathae; Passeriformes; Muscicapidae; Ficedula.
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Diadema mexicanum.
Mitochondrinon.
Eukaryota; Metazoa; Echinodermata; Eleutherozoa; Echinozoa;
Echinoidea; Euechinoidea; Diadematacea; Diadematoida; Diadematidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE FROM N.A.
STRAIN=CC70, and CC117;
MEDLINE=21561594; PubMed=11703875;
Lessios H.A., Garrido M.J., Kessing B.D.;
"Demographic history of Diadema antillarum, a keystone herbivore on Caribbean reefs.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE FROM N.A.

STRAIN=BC5, and BC8;

MEDLINE=21918460; PubMed=11918793;

Primmer C.R., Borge T., Lindell J., Saetre G.-P.;

"Single-nucleotide polymorphism characterization in species with limited available sequence information: high nucleotide diversity revealed in the avian genome.";

MOI. Ecol. 11:603-612(2002).

EMBL; AF454217; AAM22903.1; -.

EMBL; AF454218; AAM22904.1; -.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match

30.2%; Score 19; DB 8; Length 9;
Best Local Similarity 100.0%; Pred. No. 8.3e+05;
Matches 2; Conservative 0; Mismatches 0; Indels
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EMBL, AY012920; AAL33843.1; -.
EMBL, AY012921; AAL33844.1; -.
                                                                                                                                                                                           01-MAR-2002 (TrEMBLrel. 20, Last sequence update) 01-MAR-2002 (TrEMBLrel. 20, Last annotation update) Cytochrome oxidase subunit II (Fragment).
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NON TER
1 1 1 SEQUENCE 9 AA; 1174 MW; 2B73173B46DDC2D3 CRC64;
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01-OCT-2002 (TrEMBLrel. 22, Last sequence update)
01-OCT-2002 (TrEMBLrel. 22, Last annotation update)
Myelin proteolipid protein (Fragment).
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STRAIN-CC70, and CC117;
MEDLINE=21323357; PubMed=11430656;
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Lepidosauria; Squamata; Scleroglossa; Anguimorpha; Varanidae; Varanus.
NCBI_TaxID=169851;
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Mitochondrial DNA evidence and evolution in Varanoidea (Squamata).";
(Indistics 17:0-0(2001).
EMBL; AF407521; AAL10116.1; -.
Mitochondrion.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Archosauria; Aves; Neognathae; Passeriformes; Muscicapidae; Ficedula.
NCBI_TaxID=46689;
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Primmer C.R., Borge T., Lindell J., Saetre G.-P.;
Primmer C.R., Borge T., Lindell J., Saetre G.-P.;
Single-uncleotide polymorphism characterization in species with limited available sequence information: high nucleotide diversity revealed in the avian genome.";
Mol. Ecol. 11:603-612(2002).
Mol. Ecol. 11:603-612(2002).
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                                                       DB 13; Length 10;
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Pred. No. 6.1e+03;
1; Mismatches 1; Indels
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1FAAC9676732C86B CRC64;
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01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
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01-OCT-2002 (TrEMBLrel. 22, Last sequence update)
01-OCT-2002 (TrEMBLrel. 22, Last annotation update)
Myelin proteolipid protein (Fragment).
Ficedula hypoleuca.
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"Analysis of products of the Escherichia coli genomic genes and regulation of their expressions: an applicable procedure for genomic analysis of other microorganisms.";

EMBL, D21143; BAA04679.1;

Hypothetical protein.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Amphibia; Batrachia; Anura; Mesobatrachia; Pipoidea; Pipidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Bacteria, Proteobacteria, Gammaproteobacteria, Enterobacteriales,
Enterobacteriaceae, Escherichia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Steelman S., Moskow J.J., Muzynski K., North C., Druck T.,
Montgomery J.C., Huebner K., Daar I.O., Buchberg A.M.;
"Identification of a conserved family of Meisl-related homeobox
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 17; DB 2; Length 10;
Pred. No. 1.4e+04;
1; Mismatches 4; Indels
                                                          Indels
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01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-JAN-1999 (TrEMBLrel. 09, Last annotation update)
Hypothetical 1.1 kDa protein (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NON TER 1 1 1 SEQÜENCE 8 AA; 1187 MW; 278B51F37B11F40B CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Last sequence update)
Last annotation update)
                              Pred. No. 8.3e+05; 
; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                  8 AA
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             100.0%; Prea. ....
                                                                                                                                                                                                                                                                                                                                                                                                                               Created)
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Xenopus laevis (African clawed frog)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRT;
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MEDLINE=94162733; PubMed=7764507;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE FROM N.A.
MEDLINE=97202105; PubMed=9049632;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             27.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Genome Res. 7:142-156(1997).
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10 AA; 1109 MW;
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Best Local Similarity 37.5%;
Matches 3; Conservative
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TRANSFAC; T03410; -.
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                                                                 Conservative
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                                  Best Local Similarity
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                                                                                                                               8 RPG 10
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                                                                                                                                                                                         RPG 7
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P79940
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Q47561
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STRAIN-Sprague-Dawley; TISSUE-Liver;
MEDLINE-20198203; PubMed=10731662;
Kim K.-Y., Choi I., Kim S.-S.;
Kim K.-Y., Choi I., Kim S.-S.;

"Purification and characterization of a novel inhibitor of the proliferation of hepatic stellate cells.";

J. Biochem. 127:23-27(2000).
-!- FUNCTION: MAY ACT AS A NEGATIVE EFFECTOR IN THE REGULATION OF THE HEPATIC STELLATE CELLS (HSC). ALSO INHIBITS THE GROWTH OF BOVINE ENDOTHELIAL CELLS AND 316 FIBROBLASTS.
-!- SMILLARITY: IDENTICAL TO THE 63-70 AA REGION OF THE RAT ZAG
                                                                                                                                                                                                                                                                                                                                                                                                           01-0CT-2000 (TrEMBLrel. 15, Created)
01-0CT-2000 (TrEMBLrel. 15, Last sequence update)
01-0CT-2000 (TrEMBLrel. 16, Last annotation update)
01-MAR-2001 (TrEMBLrel. 16, Last annotation update)
38kDa non-arginase growth inhibitory factor (NAGIF) (Fragment).
Rattus norvegicus (Rat).
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Rattus.
NCBI_TaxID=10116;
                                                                 Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 11; Length 8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 28.6%; Score 18; DB 11; Length 8; Best Local Similarity 75.0%; Pred. No. 8.3e+05; Matches 3; Conservative 0; Mismatches 1; Indels
                                                                 Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Viruses, dsDNA viruses, no RNA stage; Herpesviridae;
Alphaherpesvirinae; Simplexvirus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    01-NOV-1996 (TrEMBLrel. 01, Last sequence update) N1-MAR-2002 (TrEMBLrel. 20, Last annotation update) mmediate-early transactivator 110 (Fragment).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        8 8 AA; 914 MW; 80A3676B02D76B1D CRC64;
Pred. No. 8.3e+05;
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                                                                 1; Mismatches
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                                     66.7%;
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                                                                 2; Conservative
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                                     Best Local Similarity
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P82598
                                                                 Matches
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                                                                                                                                                                                                       SEQUENCE FROM N.A.
STRAIN=62A, and NCTC 2916;
Dineen S.S., Bradshaw M., Johnson E.A.;
Comparison of the neurotoxin gene clusters in Clostridium botulinum type A strains.";
Submitted (DEC-2001) to the EMBL/GenBank/DDBJ databases.
EMBL; AF461539; AAM75954.1;
EMBL; AF461541; AAM75962.1;
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01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-NOV-1998 (TrEMBLrel. 08, Last annotation update)
01-NOV-1998 (TrEMBLrel. 08, Last annotation update)
Albumin (Fragment).
Aegilops squarrosa.
Bukaryota, Viridiplantae, Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; Pooideae;
Triticeae, Aegilops.
NCBL_TaxID=37682;
                                                01-0CT-2002 (TrEMBLrel. 22, Created)
01-0CT-2002 (TrEMBLrel. 22, Last sequence update)
01-0CT-2002 (TrEMBLrel. 22, Last annotation update)
BONT/A (Fragment).
BONT/A. (Fragment).
Clostridium botulinum.
Clostridium botulinum.
Bacteria; Firmicutes; Clostridia; Clostridiales; Clostridium.
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Seed storage protein.
NON TER 10 10 10 SEQÜENCE 10 AA; 1105 MW; 3A1AB5AEA365A367 CRC64;
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Pred. No. 1.4e+04;
1; Mismatches 1; Indels
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SEQUENCE 10 AA; 1143 MW; 8721FA0B1863787A CRC64;
                               10 AA.
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Job time : 35 secs
                               PRT;
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60.0%;
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Best Local Similarity 60.0 Matches 3; Conservative
                             PRELIMINARY;
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                                                                                                                                                                                 NCBI_TaxID=1491;
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Q99213
RESULT 14
Q8KHN9
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OM protein

Run on:

Sequence:

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Sequence 9, Application US/07714540
Patent NO. 5562521
GENERAL INFORMATION:
APPLICANT: Almquist, Ronald G.
APPLICANT: Toll, Lawrence
TITLE OF INVENTION: ISOLATED ATRIAL PEPTIDE-DEGRADING
TITLE OF INVENTION: ENZYME AND NOVEL COMPOUNDS USEFUL AS INHIBITORS THEREOF NUMBER OF SEQUENCES: 13
CORRESPONDENCE ADDRESSS:
ADDRESSEE: ITELL & Manella
STREET: 545 Middlefield Road, Suite 200
CITY: Menlo Park
STATE: California
             Sequence 18, Appl
Sequence 77, Appl
Patent No. 5168061
Patent No. 5169865
Patent No. 5488036
Patent No. 5488036
Patent No. 5492893
Sequence 2, Appli
Sequence 6, Appli
Sequence 6, Appli
                                                                                                                                            2, Appli
4, Appli
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14, Appl
16, Appl
19, Appl
2, Appli
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Sequence 19,
Sequence 2, A
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Sequence 12,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM FC compatible
OPERATUS SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/714,540
FILING DATE: 19910607
CLASSIFICATION: 530
                                                 5168061-1
516865-10
516985-10
5480935-1
5492893-1
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US-09-639-483C-3
US-09-383-912-18
US-09-770-014-77
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NAME: Reed, Dianne E.
REGISTRATION NUMBER: 31,292
REFERENCE/DOCKET NUMBER: 8500-
TELEPONE: 415-327-7250
TELEPAN: 415-327-7251
TELESTAN: 615-327-7251
TELESTAN: 616-327-7251
TELESTAN: 60141
INFORMATION FOR SEQ ID NO: 9:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 10 amino acids TYPE: AMINO ACID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 100.
Best Local Similarity 100.
Matches 10; Conservative
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US-07-714-540-9
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Sequence 3, Appli
Sequence 5, Appli
Sequence 22, Appli
Sequence 22, Appli
Sequence 11, Appli
Sequence 1, Appli
Sequence 6, Appli
Sequence 1, Appli
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                                                                                                                      November 17, 2003, 18:19:28; Search time 21 Seconds (without alignments) 20:148 Million cell updates/sec
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Sequence
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1: /cgn2_6/ptodata/1/iaa/5A_COMB.pep:*
2: /cgn2_6/ptodata/1/iaa/6A_COMB.pep:*
3: /cgn2_6/ptodata/1/iaa/6A_COMB.pep:*
4: /cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*
5: /cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*
5: /cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*
                   GenCore version 5.1.6
(c) 1993 - 2003 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-08-343-883-1

US-08-342-883-1

US-08-341-219-11

US-08-453-588-22

US-08-453-588-22

US-08-459-917-1

US-08-465-18

US-08-64-865-18

US-08-486-18

US-08-481-18

US-08-471-1758-6

US-08-21-079-22

US-09-100-4148-77

US-09-100-4148-77

US-09-91-23-128-13
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US-07-690-983D-32
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-09-698-134-1
US-09-256-599-1
                                                                                                                                                                                                                                                                                                                                  328717 segs, 42310858 residues
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Maximum Match 100%
Listing first 45 summaries

    protein search, using sw model

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Gapop 10.0 , Gapext 0.5
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63
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Match Length
                                  Copyright
                                                                                                                                                                                                                                                                                                                                                                                                         seq length: 0
seq length: 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            litle:
Perfect score:
                                                                                                                                                                                                                                                                              Scoring table:
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Minimum DB Maximum DB

Database

Result

Searched:

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APPLICANT: Dufour, Raymond J.
APPLICANT: Roulet, Claude J.M.
APPLICANT: Roulet, Claire D.
APPLICANT: Bonneau, Michel B.
TITLE OF INVENTION: Method for improving the organoleptic
TITLE OF INVENTION: qualities of the meat from uncastrated male domestic
TITLE OF INVENTION: animals, vaccines which are usable in this method, new
TITLE OF INVENTION: peptide, in particular for producing these vaccines...
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Indels '0;
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/343,883
FILING DATE: 17-NOV-1994
PIOR APPLICATION NUMBER: 08/08/343,883
FILING DATE: 07/0944
PRIOR APPLICATION NUMBER: 08/0946,495
FILING DATE: 09-NOV-1992
PRIOR APPLICATION NUMBER: RF 9102513
FILING DATE: 01-MAR-1991
SOFTWARES PATENTIN RALESSE #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/690,983D
FILING DATE: 25-JUN-1991
CLASSIFICATION DATE: PS-JUN-1991
CLASSIFICATION DATE: PCT/AU90/00373
APPLICATION NUMBER: PCT/AU90/00373
FILING DATE: 24-AUG-1990
ATTONREY/AGENT INPOMBER: 29,768
REGISTRATION NUMBER: 29,768
REGISTRATION NUMBER: 16786/148 CHAC
TELECOMMUNICATION: INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202)672-5300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        E: Larson and Taylor
727 Twenty-Third Street, South
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ZIE: 22202
COMPUTER READABLE FORM:
MEDIUM TYPE: Flopy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE: 01-MAR-1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: FF 9115289
FILING DATE: 10-DEC-1991
INFORMATION FOR SEQ ID NO: 1:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; Sequence 1, Application US/08343883
; Patent No. 5573767
                                                                                                                                                                                                                                                                                                                                    TELEFAX: (202) 672-5399
INFORMATION FOR SEQ ID NO: 32:
SEQUENCE CHARACTERISTICS:
LENGTH: 10 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Best Local Similarity 100.
Matches 10; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MOLECULE TYPE: protein
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SY: linear
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STATE: Virginia
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY:
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US-08-343-883-1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/O7/690,983D
FILING DATE: 25-JUN-1991
CLASSIFICATION: 435
RIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/AU90/00373
FILING DATE: 24-AUG-1990
ATTORNEY/AGENT INFORMATION:
NAME: BENT, Stephen A.
REGISTRATION NUMBER: 29,768
REFERENCE/DOCKET NUMBER: 16786/148 CHAC
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0; Mismatches
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US-07-690-983D-32
US-07-690-983D-32
Sequence 32, Application US/07690983D
Patent No. 503586
GENERAL INFORMATION:
APPLICANT: RUSSELL-JONES, Gregory J. APPLICANT: STEWART, Andrew G. APPLICANT: TSONIS, Con G. TITLE OF INVENTION: FUSION PROTEINS NUMBER OF SEQUENCES:
CORRESPONDENCE ADDRESS:
ADDRESSEE: FOLEY & Lardner
                                                                                                                                              APPLICANT: RUSSELL-JONES, Gregory J. APPLICANT: STEWART, Andrew G. TPPLICANT: TSONIS, Con G. TITLE OF INVENTION: FUSION PROTEINS NUMBER OF SEQUENCES: 47
ADDRESSEE: FOLEY & Lardner
                                                                                                                                                                                                                                                                                                                                                                             ZIP: 20007-5109
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBB PC compatible
OPERATING SYSTEM: PC-DOS/MS_DOS
                                                                      Sequence 2, Application US/07690983D Patent No. 5403586 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STREET: Boley & Lardner STREET: 3000 K Street, N.W. CITY: Washington, D.C. COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ZIP: 20007-5109
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                  STREET: FOLGY & Lardner STREET: 3000 K Street, N.W. CITY: Washington, D.C. COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEPHONE: (202)672-5300
TELEPAX: (202)672-5399
INFORMATION FOR SEQ ID NO: 2: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    : 10 amino acids
amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 100.
Best Local Similarity 100.
Matches 10; Conservative
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EHWSYGLRPG 10
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                                                          US-07-690-983D-2
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Sequence 22, Application US/08428488
Sequence 22, Application US/08428488
Patent No. 5624991
GENERAL INFORMATION:
APPLICANT: BODOR, Nicholas S,
TITLE OF INVENTION: PERTIDES BY SEQUENTIAL METABOLISM
NUMBER OF SEQUENCES: 107
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 100.0%; Score 63; DB 1; Length 10; Best Local Similarity 100.0%; Pred. No. 0.00011; Matches 10; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                           Query Match 100.0%; Score 63; DB 1; Length 10; Best Local Similarity 100.0%; Pred. No. 0.00011; Matches 10; Conservative 0; Mismatches 0; Indels
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US-08-428-488-22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STATE: Algorithm of the States
COUNTRY: United States
ZID: 22313-1404
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/428,488
FILING BAYE: 27-APR-1995
CLASSIFICATION: 514
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OTHER INFORMATION: /note= "Position 1 = p-Glu."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE: Burns, Doane, Swecker & Mathis STREET: P.O. Box 1404
CITY: Alexandria
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ATTORNEY/AGENT INFORMATION:
NAME: Baumeister, Mary Katherine
REGISTRATION NUMBER: 26,254
REFERENCE/DOCKET NUMBER: 028724-087
TELECOMMUNICATION INFORMATION:
TELEPAX: (703) 83-6620
TELEFAX: (703) 83-2021
INFORMATION FOR SEQ ID NO: 22:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME/KEY: Modified-site
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                    SEQUENCE CHARACTERISTICS:
LENGTH: 10 amino acids
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LENGTH: 10 amino acids
    INFORMATION FOR SEQ ID NO:
                                                                                        single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                        TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                            1 EHWSYGLRPG 10
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                                                                                                                                                                                                                                                                                      1 EHWSYGLRPG 10
                                                               TYPE: amino acid
STRANDEDNESS: sir
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Patent No. 5578477
GENERAL INFORMATION:
PAPLICANT: Tamanoi Dr., Fuyuhiko
TITLE OF INVENTION: IDENTIFICATION AND CHARACTERIZATION OF
TITLE OF INVENTION: INHIBITORS OF PROTEIN FARNESYLTRANSFERASE
NUMBER OF SEQUENCES: 10
CORRESPONDENCE ADDRESS:
ADDRESSEE: Foley & Lardner
STREET: 3000 K Street, N.W., Suite 500
COUNTRY: Washington, D.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    .
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Structure of the porcine LH- and FSH-releasing hormone. I. The proposed amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      100.0%; Score 63; DB 1; Length 10; 100.0%; Pred. No. 0.00011; tive 0; Mismatches 0; Indels
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ZIP: 2000-5109

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/000,931
FILING DATE: O-JAN-1994
CLASSIPICATION: 435
ATTONNEY/AGENT INPORMATION:
NAME: BENT, Stephen A
REGISTRATION NUMBER: 29,768
REGISTRATION NUMBER: 29,768
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEFRAX: (202)672-5309
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ) DATE: 1971

) RELEVANT RESIDUES IN SEQ ID NO: 1: FROM 1 TO 10

US-08-343-883-1
                                                                                                                                                                                                                  FEATURE:
NAME/KEY: Peptide
LOCATION: 1
OTHER INFORMATION: /label= pyro
OTHER INFORMATION: /note= "pyroglutamic acid"
AUTHORS: Matsuo, H.O.
                                                                                                                                                     LOCATION: 10 OTHER INFORMATION: /label= NH2 OTHER INFORMATION: /note= "amidated glycine"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Biochem. Biophys. Res. Commun.
    LENGTH: 10 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: peptide
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                       Baba, Y.
G. Nair, R. M.
Arimura, A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity 100.
Matches 10; Conservative
SEQUENCE CHARACTERISTICS
LENGTH: 10 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                       Schally, A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 EHWSYGLRPG 10
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                                                                                                                                NAME/KEY: Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1334-1339
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TELEX: 904136
                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE: St
TITLE: FS
TITLE: se
JOURNAL:
                                                                                                                                                                                                                                                                                                                                                                       AUTHORS:
AUTHORS:
AUTHORS:
AUTHORS:
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ISSUE: 6
PAGES: 1
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Sequence 22, Application US/08453588
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                                                                                                                                                APPLICANT: Zohar, Y.
APPLICANT: Rivier, J.
APPLICANT: Rowell, J.
APPLICANT: Schaul, J.
APPLICANT: Gordilf, Y.
TITLE OF INVENTION: Compounds and Methods For Controlling
TITLE OF SEQUENCES: 26
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                100.0%; Score 63; DB 1; Length 10; 100.0%; Pred. No. 0.00011; cive 0; Mismatches 0; Indels
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CURRENT APPLICATION DATA:

APPLICATION NUMBER:

FLING DATE: 05/08/341,219

FLING DATE: 05/08/341,219

CLASSIFICATION: 514

ATTONREY/AGENT INPORMATION:

NAME: CCLUZZi, Laura A.

REGISTRATION NUMBER: 30742

REFERENCE/POCKET NUMBER: 399-003-999

TELEPHONICATION INFORMATION:

TELEPHONE: (212) 790-9090

TELEPHONE: (212) 790-9090
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /note= "=pyroglutamic acid"
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/label= Gly10
/note= "amidated"
                                                                                                                                                                                                                                                                                                                                              ADDRESSE: Pennie & Edmonds
STREET: 1155 Avenue of the Americas
CIIY: New York
STATE: N.Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /product= "OTHER'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                            Sequence 11, Application US/08341219
Patent No. 5643877
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: amino acid
STRANDEDNESS: not relevant
TOPOLOGY: unknown
MOLECULE TYPE: peptide
HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                      ZIP: 10036-2711
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FEATURE:
NAME/KEY: Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME/KEY: Modified-site
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OTHER INFORMATION: /
OTHER INFORMATION: /
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EHWSYGLRPG 10
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  1 EHWSYGLRPG 10
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OTHER INFORMATION:
OTHER INFORMATION:
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   HYPOTHETICAL: NC
ANTI-SENSE: NO
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                USA
                                                                               US-08-341-219-11
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RESULT 8 US-08-453-588-22

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Gaps
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                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

CURRENT APPLICATION DATA:

APPLICATION NUMBER: 08/078,661

PILING DATE: 16-JUN-1993

ATTORNEY/AGENT INFORMATION:

NAME: MATY E. GOTTHLEY

REGISTRATION NUMBER: 34,49

TELECOMMUNICATION INFORMATION:

NAME: MATY E. GOTTHLEY

REGISTRATION NUMBER: 34,49

TELECOMMUNICATION INFORMATION:

TELECOMMUNICATION INFORMATION:

REGISTRATION POR SEQ ID NO: 22:

SEQUENCE CHARACTERISTICS:

LUMBER: 10 aming acide
Patent No. 568415
GENERAL INFORMATION:
APPLICANT: Anna van der Zee, Irma Marianne van Die,
APPLICANT: Willem Pieter Martin Hoekstra,
APPLICANT: Josephus Theodorus Gielen.
TITLE OF INVENTION: Carrier system against GnRH
NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: Denver
STATE: Colorado
CUNTRY: USA
ZIP: 80203
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME/KEY: Glu at position 1 is pyroglutamic acid
                                                                                                                                                                                                              ADDRESSE: Akzo No. 5684145el Patent Department STRET: 1300 Piccard Drive, Sulte 206 CITY: Rockville STATE: Maryland
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 1, Application US/08591917
Patent No. 5707964
GENERAL INFORMATION:
APPLICANT: Nett, Torrance M
APPLICANT: Glode, Leonard Michael
TITLE OF INVENTION: A METHOD FOR TREATING CANCER
NUMBER OF SEQUENCES:
ADDRESSEE: Sheridan Ross & McIntosh
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
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STREET: 1700 Lincoln Street, Suite 3500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 .,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity 100.
Matches 10; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MOLECULE TYPE: protein
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Gaps
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Parent No. 5827659
GENERAL INFORMATION:
APPLICANT: PATTERSON, DALE H.
APPLICANT: TARK GEORGE E.
TITLE OF INVENTION: METHODS AND APPARATUS FOR SEQUENCING TITLE OF INVENTION: POLYMERS USING MASS SPECTROMETRY.
NUMBER OF SEQUENCES:
ADDRESSEE: Patent Administrator - Testa, Hurwitz & ADDRESSEE: Thibeault
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 10;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: Floppy disk
COMPUTER: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/796,598
FLING DATE: 07-FEB-1997
CLASSIFICATION DATE: WAY-1997
PRIOR APPLICATION DATE: US/08/796,598
FLING DATE: 19-MAY-1995
ATTORNEY/AGENT INFORMATION:
NAME: FLYNN ESQ., KEATY A.
REGISTRATION NUMBER: 33,693
REFERENCE/DOCKET NUMBER: SYP-115
TELECOMMUTICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%; Score 63; DB 2; I 100.0%; Pred. No. 0.00011;
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Sequence 18, Application US/08694865
Sequence 18, Application US/08694865
GENERAL INFORMATION:
APPLICANT: POTTER, ANDREW A.
APPLICANT: MANNS, JOHN G.
TITLE OF INVENTION: GRH-LEUKOTOXIN CHIMERAS
NUMBER OF SEQUENCES: 34
CORRESPONDENCE ADDRESS:
ADDRESSER: REED & ROBINS LLP
STREET: SES HAMILTON AVENUE, SUITE 200
CITY: PALO ALTO
                                                                                                                                                                                                                                                                                                                                                                                                                                          STREET: High Street Tower, 125 High Street CITY: Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
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INFORMATION FOR SEQ ID NO: 6
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 10 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity 100.
Matches 10; Conservative
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy o
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         , MOLECULE TYPE: peptide US-08-796-598-6
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                                           1 EHWSYGLRPG 10
                                                                       1 EHWSYGLRPG 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY:
                                                                                                                                                                            US-08-796-598-6
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APPLICANT: Ladd, Anna
APPLICANT: Ladd, Anna
APPLICANT: Ladd, Anna
APPLICANT: Lang, Vi
APPLICANT: Lamb, Timothy
TITLE OF INVENTION: Immunogenic LHRH peptide constructs
TITLE OF INVENTION: and synthetic universal immune stimulators for vaccines
NUMBER OF SEQUENCES: 114
CORRESPONDENCE ADDRESS:
ADDRESSEE: Maria C.H. Lin
STREET: 345 Park Avenue
CITY: New York
STATE: NY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%; Score 63; DB 1; Length 10; 100.0%; Pred. No. 0.00011; Live 0; Mismatches 0; Indels
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ZIP: 10154-0053

ZIP: 10154-0053

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/446,692
FILING DATE: 7-UNN-1995
CLASSIFICATION: 424
ATYONEY/AGENT INFORMATION:
NAME: MAIA C.H. Lin REGISTRATION NUMBER: 29,323
RESPENCE/DOCKET NUMBER: 29,323
RESPENCE/DOCKET NUMBER: 1151-4146 US2
TELECOMMUNICATION INFORMATION:
TELEPRAK: (516)751-6849
                                                                                                      ATTORNEY/AGENT INFORMATION:
NAME: Kovarik, Joseph E.
REGISTRATION NUMBER: 33,005
REFERENCE/DOCKET NUMBER: 2730-3-2-1-1
TELECOMMUNICATION INFORMATION:
TELEPHONE: (303) 863-9700
TELEPAX: (303) 863-9700
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 10 amino acids
26-JAN-1996
26-JAN-1996
NN: 514
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 1, Application US/08446692
Patent No. 5759551
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 10 amino acids
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; TOPOLOGY: linear;
; MOLECULE TYPE: peptide
US-08-446-692-1
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Best Local Similarity 100.8
Matches 10; Conservative
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Best Local Similarity 100.
Matches 10; Conservative
                                                                                                                                                                                                                                                                                                                                     TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
US-08-591-917-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 EHWSYGLRPG 10
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                                        APPLICATION NUMBER:
FILING DATE: 26-JAN
CLASSIFICATION: 514
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US-08-446-692-1
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Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                100.0%; Score 63; DB 2; Length 10; 100.0%; Pred. No. 0.00011;
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STATE: Massachusetts
CUNTRY: USA
ZIP: 02109-1875
COMPUTER READABLE FORM:
MEDTUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATIOS SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATONIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/480,494B
FILING DATE: 07-JUN-1995
CLASSIFICATION: 514
ATYONEN'APARTI INFORMATION:
NAME: Deconti, Giulio A.
REFERENCE/DOCKET NUMBER: 31,503
REFERE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 1, Application US/08480494B
Patent No. 5843901
GENERAL INFORMATION:
GENERAL INFORMATION:
TITLE OF INVENTION: LHRH Antagonist Peptides
NUMBER OF SEQUENCES:
CORRESPONDENCE S:
ADDRESSEE: LAHIVE & COCKFIELD
STREET: 60 State Street, Suite 510
CLASSIPTCATION: 424

PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/057,166

FILING DATE: 27-APR-1992

CLASSIFICATION: 424
ATTORNEY/ABONT INFORMATION:
NAME: Maria C.H. Lin
RECISTRATION NUMBER: 29,323

REFERENCE/DOCKET NUMBER: 1151-4146 US2
TELECHOMNICATION INFORMATION:
TELECHOME: (212/415-849)
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 10 main o acids
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Best Local Similarity 100.
Matches 10; Conservative
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MOLECULE TYPE: peptide
US-08-488-351A-1
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APPLICANT: Wang, Timchu
TITLE OF INVENTION: Immunogenic Lithi peptide constructs
TITLE OF INVENTION: and synthetic universal immune stimulators for vaccines
TITLE OF INVENTION: and synthetic universal immune stimulators for vaccines
TITLE OF INVENTION: 114
CORRESSEE: Maria C.H. Lin
STREET: 345 Park Avenue
CITY: New York
STRTE: Ny
COMPUTE: Ny
COUNTY: New York
STATE: NY
COMPUTER: Ploppy disk
COMPUTER: Ploppy disk
COMPUTER: Ploppy disk
COMPUTER: Ploppy disk
COMPUTER: PLOPS/MS-DOS
SOFTWARE: Petentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
PRILING DATE: 7-UUM-1995
CLASSIFICATION DATA:
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
TILING DATE: 7-UUM-1995
CLASSIFICATION: 424
PRIOR APPLICATION: 424
PRIOR APPLICATION: 424
PRIOR APPLICATION: 424
PRIOR APPLICATION: 424
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Best Local Similarity 100.0%; Pred. No. 0.00011;
Matches 10; Conservative 0; Mismatches 0; Indels
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CTHER INFORMATION: /note= "This position is pyroGlu."
US-08-694-865-18
          COMPUTER READBLE FORM:
MEDDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/694,865
FILING DATE: 09-AUG-1996
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: MCCRACKEN, THOMAS P.
REFERENCE/DOCKET NUMBER: 38,548
REFERENCE/OFFENTION INFORMATION:
TELEPHONE: (415)327-3231
INFORMATION POR SEQ ID NO: 18:
SEQUENCE CHARACTERISTICS:
LENGTH: 10 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 1, Application US/08488351A Patent No. 5843446 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME/KEY: Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY: linear
MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 EHWSYGLRPG 10
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EHWSYGLRPG 10

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US-00-474-175A-6

US-00-474-175A

US-00-474-175A

Patent No. 586940

EARLAL INCRATION

TITLE OF INVENTION: METHODS AND APPARATUS FOR SEQUENCING

TITLE OF INVENTION: SPECTROMETRY.

NUMBER OF SEQUENCES: 23

CORRESPONDENCE ADDRESS: 23

CORRESPONDENCE ADDRESS: 34

COURREY DESCRIPTION: DIA SPECTROMETRY.

ADDRESSEE: Patent Administrator - Testa, Hurwitz & COUNTRY: USA

ITTLE OF INVENTION: SPECTROMETRY: USA

ITTLE OF INVENTION: DIA STREET: High Street Tower, 125 High Street

COUNTRY: USA

ITTLE OF INVENTION: DIA STREET: HIGH STREET This Dead the Compatible

COUNTRY: USA

INFORMATION OF SECONDERS: SOFTWARE: Patentin Release #1.0, Version #1.30

COMPUTER: ISA PECONDERS: SOFTWARE: Patentin Release #1.0, Version #1.30

CLASSIFICATION NUMBER: US/08/447,175A

FILING DATE: 19-MAY-1995

CLASSIFICATION NUMBER: SYP-114

TELECHOMET ARDRESSITION: TOWERS: SYP-114

TELECHOMET (617) 248-7000

TELECHOMET (617) 248-7000

TELECHOMET OF SECONDERS STREET

TYPE: Amino acids

TYPE: Amino acids
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Best Local Similarity 100.
Matches 10; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: peptide
US-08-447-175A-6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 EHWSYGLRPG 10
1 EHWSYGLRPG 10
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Search completed: November 17, 2003, 18:23:37 Job time : 21 secs

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protein

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Run on:

Sequence:

Searched:

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Lutenizing hormone
Rat GnHR peptide.
Neutrophil-activat
Ubiquitin fusion p
Non-crosslinked pr
                                                                                                                                                                  Luteinising hormon
Gonadorelin peptid
Human LHRH peptide
Gonadotropin relea
Luteinising hormon
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LHRH target antige
Luteinising hormon
Native mammalian g
                           Gonadotropin relea
Gonadotropin relea
Luteinising hormon
Luteinising hormon
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Gonadorelin peptid
Gonadotropin relea
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Luteinising hormon
Hormone domain of
                                                                                                                                                                                                                              Amino acid sequenc
Luteinising hormon
Gonadotrophin rele
                                                                                                                                                                                                                                                           Gonadotrophin-Rele
Mammalian releasin
releasing hormo
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                                                                   Luteinising
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LHRH; Follicle Stimulating Factor; FSH; acne; hirsutism; dysmenorrhea; precocious puberty; endometriosis; prostate cancer; benign prostate hypertrophy; mammary tumour.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      acid"
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                                                                  AAW45642
AAW04612
AAW76373
AAY5022
AAY31176
AAY31067
AAY3856
AAY3866
AAW96765
AAW96765
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AAB08103
AAB03590
AAB06261
AAY88576
AAY79054
AAY91197
AAY68566
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AAB10930
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AAB20863
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 83BE-0468932
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
 /note=
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(updated)
(updated)
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Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    23-FEB-1983;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       25-MAR-2003
10-MAR-2003
01-JUL-2002
17-DEC-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Luteinising
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 AAP10411;
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Luteinising hormon
Sequence of lutein
Peptide #1 with ho
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Luteinising Hormon
Gonadotrophin rele
Gonadoliberin anta
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Luteinising Hormon
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Novel decapeptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                               November 17, 2003, 18:06:32 ; Search time 41 Seconds (without alignments) 38.714 Million cell updates/sec
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| SID
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        GenCore version 5.1.6
Copyright (c) 1993 - 2003 Compugen Ltd.
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Maximum Match 100%
Listing first 45 summaries
                                               sw model
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AAP10416
AAP50222
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AAP70922
AAP90630
AAR15713
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Gapop 10.0 , Gapext 0.5
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Minimum I Maximum I

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Result Š

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The claimed peptide antagonists inhibit the release of gonadotrophins and/or steroids. They are antagonistic to {\tt GNRH}, inhibit ovulation, and
                                                                                                                                                                                  A composition is claimed containing LHRH or its analogues. The composition is used to treat prostate adenocarcinoma, benign hypertrophy of the prostate, endometriosis, dysmenorrhea, hirsutism, hormone-dependent mammary tumours, for treatment or prevention of precocious puberty, delaying the onset of puberty and for treating acne. The compositions may also contain antiandrogens.
                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                       and its agonists compsn.
benign hypertrophy of
                                                                                                                                                                                                                                                                                                                                                                                             .
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100.0%; Pred. No. 0.00032;
.ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GnRH; LH-RH; LRF; gonadotrophins; steroids; contraceptive.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New GnRH antagonist peptide(s) - useful as inhibitors of gonadotropin(s) and/or steroid(s) for contraceptive use.
                                                                                                                                                                                                                                                                  See APP10411-P10418
(Updated on 01-JUL-2002 to add missing PI field.)
(Updated on 10-MAR-2003 to add missing OS field.)
(Updated on 25-MAR-2003 to correct PA field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gonadotrophin release stimulating hormone.
                                                                                                       LH-RH, liberating factor for LH and FSH, - used to treat prostate adenocarcinoma, the prostate, hirsutism, acne, etc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (SALK ) SALK INST BIOLOGICAL STUDIES.
                                                                                                                                                             Claim 1(f); Page 16; 27pp; French.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAP50222 standard; Protein; 10 AA.
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                                                                             WPI; 1981-23409D/14 (23409D)
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79FR-0023545.
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85US-0771517
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(first entry)
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Best Local Similarity 100.
Matches 10; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                          FHWSYGLRPG 10
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                          (ROUS ) ROUSSEL-UCLAF
                                                     Raynaud J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 1985-136434/23.
                                                                                                                                                                                                                                                                                                                                         10 AA;
 21-SEP-1979;
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20-JAN-1992
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                                                                                                                                                                                                                                                                                                                                           Sequence
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                                                     Labrie F,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic.
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                                                                                                                                                                                      A composition is claimed containing LHRH or its analogues. The composition is used to treat prostate adenocarcinoma, benign hypertrophy of the prostate, endometriosis, dysmenorrhea, hirsutism, hormone-dependent mammary tumours, for treatment or prevention of precocious puberty, delaying the onset of puberty and for treating acne. The compositions may also contain antiandrogens.

See also AAR10412-P10418.
(Updated on 01-JUL-2002 to add missing PI field.)
(Updated on 10-MAR-2003 to add missing OS field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LHRH; Follicle Stimulating Factor; FSH; acne; hirsutism; dysmenorrhea; precocious puberty; endometriosis; prostate cancer; benign prostate hypertrophy; mammary tumour.
                                                                                                       LH-RH, liberating factor for LH and FSH, and its agonists compsn. - used to treat prostate adenocarcinoma, benign hypertrophy of the prostate, hirsutism, acne, etc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /note= "amidated or absent, in which case Pro(9)
is Pro-NH-C2H5"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             'note= "N-alpha-methyl-Leu"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /label= OTHER
/note= "pyroglutamic acid"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAP10416 standard; peptide; 10 AA
                                                                                                                                                               Claim 1(a); Page 15; 27pp; French
                                                                              WPI; 1981-23409D/14 (23409D).
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EHWSYGLRPG 10
                          (ROUS ) ROUSSEL-UCLAF
                                                     Labrie F, Raynaud J;
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tes 10; Conserv
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Modified-site
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  21-SEP-1979;
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17-DEC-1992
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AAP61403 standard; protein; 10 AA.
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                                      25-MAR-2003
09-JAN-2003
04-AUG-1991
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                                                                                                                                                                                                      08-MAY-1984;
                                                                                                                            Unidentified
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27-OCT-1991
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                    AAP61403;
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may cause resorption of a fertilised egg if administered shortly after absorption. The peptides also have utility in male contraception, and in treatment of precoscious puberty, hormone dependent neoplasia, dysmenorrhoea and endometriosis.
                                                                                              Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The decapeptide encodes a gonadoliberin antagonist, which may be used as a male contraceptive and as an antitumour (against steroid-dependent tumours).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New peptide gonadotropin releasing hormone antagonists - useful esp. as contraceptives, for treating early puberty, hormone-dependent neoplasms etc.
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                                                                            Length 10;
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                                                                          Query Match . 100.0%; Score 63; DB 6; Length 10
Best Local Similarity 100.0%; Pred. No. 0.00032;
Matches 10; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                 Gonadoliberin antagonist; contraceptive; antitumor.
                                                                                                                                                                                                                                                                                                                                                                                                                     Rivier JEF, Varga JI, Hagler AT, Struthers RS,
Rivier CL, Vale WW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (Updated on 31-OCT-2002 to add missing OS field.) (Updated on 25-MAR-2003 to correct PA field.)
                                      (Updated on 25-MAR-2003 to correct PA field.)
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                                                                                                                                                                                  AAP60127 standard; Peptide; 10 AA.
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(updated)
(first entry)
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Best Local Similarity 100.
                                                                                                                                                                                                                                                              Gonadoliberin antagonist
                                                                                                                            1 EHWSYGLRPG 10
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                                                        10 AA
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                                                                                                                                                                                                                                                                                                     Unidentified
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31-OCT-2002
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                                                                                                                                                                                                     AAP60127;
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RESULT 5 AAP61403

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The gonadotropin releasing hormone and its analogues are prepd. by a new multistage rapid peptide synthesis method in soln., where the intermediates are not isolated. The process is rapid and gives very pure peptide quickly and using little equipment. The peptide can be used in veterinary medicine to synchronise ovulation in large animal herds, and in human medicine in the treatment of fertility disorders. (Updated on 09-JAN-2003 to odd missing OS field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Prepn. of gonadotropin liberating hormone and analogues - by multistage rapid peptide synthesis in soln. without isolating
                                                                                                                          Gonadotropin releasing hormone; analogue; peptide synthesis; ovulation; veterinary medicine; fertility;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 10;
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100.0%; Pred. No. 0.00032;
iive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel decapeptide with LHRH inhibition activity.
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                                                                                                                                                                                                                                                                                                                                                                                                         (DEAK ) AKAD WISSENSCHAFTEN DDR.
                                                                                  Gonadotropin releasing hormone.
                                                                                                                                                                                                                                                                                                                                                                 84DD-0262804.
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(updated)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 100.
Best Local Similarity 100.
Matches 10; Conservative
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EHWSYGLRPG 10
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Analogues (I) of the sequence pref. have amino acid (AA) substitutions at posns. 2 (his is replaced by a D-AA) and 6 (gly is replaced by a D-AA).

A therapeutically effective amt. of such an antagonist is contained in a pharmaceutical compsn. alongside a menopausal-symtom-alleviating and. of a 19-nor progestational agent (II) (pref. both in single formulation).

The compsn. is pref. administered nasally in dosages of 0.01-1 mg/kg/ day for (I) and 0.02-0.07 mg/kg/day for (II). May be used for inhibition of ovulation, and treatment of eg endometriosis, breast cancer, polycistic ovarian disease, or precocious puberry in female mammals.

(Updated on 03-OCT-2002 to add missing OS field.)
                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Compsn. comprising LHRH-antagonist and 19-nor progestational agent for treating female gynaecological disorders based on gonads steroid producn.
This luteinising hormone releasing hormone (LHRH) agonist has either a ketomethylene or dihydroketomethylene gp. replacing the amide linkage between residues 9 and 10 in LHRH. This results in an increase in oral activity. It is useful for eg male and female contraception, treatment of precocious puberty and endometriosis and treatment of breast- and prostate tumours. See also AAP70923-27.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Luteinizing hormone releasing hormone (LHRH); LHRH antagonist; 19-nor-progestational agent; female gynaecological disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 10;
                                                                                                                                                                                   Length 10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence of luteinizing hormone releasing hormone (LHRH).
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                                                                                                                                                                                   100.0%; Score 63; DB 8; L
100.0%; Pred. No. 0.00032;
ive 0; Mismatches 0;
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les 10; Conservative
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                                                                                                                                                    10 AA;
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14-JUN-1989
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                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                            Peptide inhibits the release of lutenising hormone releasing hormone.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Orally active luteinising hormone-releasing hormone peptide analogues - have keto:methylene or hydroxy:ethylene in place of amide between proline(9) and glycine(10)
                                                                                                                                                                                                                                                                                                                                                                                                     ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /note= "other= ketomethylene(Gly), dihydroketo-
methylene(Gly)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       breast tumours; prostate tumours; ectopic tumours; menopause.
                                                                                                                                                                                                                                                                                                                                                                 Length 10;
                                                                                                                                                                                                                                                                                                                                                                                                  0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LHRH; contraception; precocious puberty; endometriosis;
                                                                                                                                                                                      Deca:peptide - inhibits LH-RH hormone release activity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /note= "other= pyroglutamic acid"
                                                                                                                                                                                                                                                                                                                                                                 100.0%; Score 63; DB 7; I 100.0%; Pred. No. 0.00032;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Luteinising hormone releasing hormone agonist
                                                                                                                                                                                                                                                                            See also AAP60575.
(Updated on 25-MAR-2003 to correct PA field.)
                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                         Disclosure; Page 990; 5pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
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                                                                                               (TULA ) ADMIN TULANE EDUCAT.
(TULA ) TULANE EDUCATIONAL FUND
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/label= other
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/label= other
                           85JP-0185616
                                                              84US-0643643
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                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                    10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       EHWSYGLRPG 10
                                                                                                                                                    WPI; 1986-321434/49.
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                                                                                                                                                                                                                                                                                                                                                                                                                                    1 EHWSYGLRPG
                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                Sequence 10 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Key
Modified-site
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                             23-AUG-1985;
                                                                23-AUG-1984;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   01-MAY-1991
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                                                                                                                                                                                                                                                                                                                                                                                                     Matches
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tumours;

RESULT 9 AAR15713

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The decapeptides is an antagonistic analogue of hypothalamic LHRH which possesses high antiovulatory and antineoplastic activity, is free of anaphylactoid side effects and is believed to be free of endematogenic effects. The peptide may be used to treat precocious prostate tumours, e.g. malignant and benign prostate tumours, e.g. secondary amenorrhoea, endometriosis and ovarian and mammary cystic diseases. The peptide is also useful for regulating ovulation e.g. as precoital or postcoital contraceptives, for synchronising oestrus in livestock and for improving the "rhythm" method. It is also useful for regulating the human menopausal and postmenopausal periods. As it suppresses
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     the spermatogenesis and testosterone levels in males, it may be of potential use for male contraception.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New deca:peptide luteinising hormone-releasing hormone antagonists - for treating precocious puberty, hormone dependent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               antagonists - for treating precocious puberty, hormone depende tumours, endometritis, cystic diseases; also as contraceptive
                                                                                                                                            Luteinising hormone; LHRH; hypothalamic; antiovulatory; tumou antineoplastic; precocious puberty; ovulation; contraceptive.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%; Score 63; DB 13;
100.0%; Pred. No. 0.00032;
ative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         See also AAR26818, AAR29046-7. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                     Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAR62689 standard; peptide; 10 AA.
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                                                                                                                                                                                                                                                                                                         /note= "amidated"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Schally AV;
                                                                                                           LH releasing hormone antagonists.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (TULA ) TULANE EDUCATIONAL FUND.
                                                                                                                                                                                                                                                                    /label= pGlu
                                                                                                                                                                                                                                                                                                                                                                                                                                                    91US-0647786.
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                                                       (updated)
(first entry)
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EHWSYGLRPG 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1992-299984/36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity
les 10; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              10 AA;
                                                                                                                                                                                                                                                    Misc-difference
                                                                                                                                                                                                                                                                                       Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                               29-JAN-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                    30-JAN-1991;
                                                       25-MAR-2003
10-FEB-1993
                                                                                                                                                                                                                                                                                                                                           WO9213883-A1
                                                                                                                                                                                                                                                                                                                                                                              20-AUG-1992.
                                                                                                                                                                                                    Synthetic.
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                      AAR26819;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
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AAR62689
ID AAR62
XX
AC AAR62
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The C-terminal residue (GJy-CO-NH2) can be replaced by ethylamide. This peptide and fragments of it (i.e. amino acids 4-10, 5-10, 6-10 and 7-10) are agonists and antagonists of LHRH. They are useful for treating e.g. precocious or delayed puberty, psychiatric disorders esp. those of the libido or sexual aggression, etc. In addition they are useful for functional exploration of the hypothalamus-hypophyseal axis and for radioimmunological to biological assay (of LH, FSH and (Updated on 25-MAR-2003 to correct PA field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New peptide homologous with luteinising hormone-releasing hormone - used to treat gynaecological conditions, cancer of gonads and sec. sexual organs, psychiatric conditions and in assays
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Bauer K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (INRM ) INSERM INST NAT SANTE & RECH MED.
                                                                                                                                                                                                                                                                                                                                           Location/Qualifiers
                                                                                                         AAR15713 standard; Protein; 10 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /label= OTHER
/note= "amidated"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Kordon C,
                                                                                                                                                                                                                                   Peptide #1 with homology to LHRH.
                                                                                                                                                                                                                                                                                                                                                                                              'note= "pyroGlu"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 3; Page 50; 83pp; French
                                                                                                                                                                                                                                                                                                                                                                          90FR-0005147.
                                                                                                                                                                                                                                                                                                                                                                                                                                 'label= Hyp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               91WO-FR00332
                                                                                                                                                                               (updated)
(first entry)
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Best Local Similarity 100.
Matches 10; Conservative
1 EHWSYGLRPG 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1991-339753/46.
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                                                                                                                                                                                                                                                                                                                                                         Modified-site
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                                                                                                                                                                             25-MAR-2003
24-JAN-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO9116343-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            31-OCT-1991.
                                                                                                                                                                                                                                                                    luliberin.
                                                                                                                                                                                                                                                                                                       Synthetic.
                                                                                                                                              AAR15713;
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Gaps

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AAR62689

AAR26819 standard; peptide; 10 AA.

RESULT 10 AAR26819

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Length 10; Indels

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Gonadotropin releasing hormone; GnRH; motility disorder; functional bowel disease; leuprolide acetate; luceinising hormone; progesterone; relaxin; autonomic nervous system; drug delivery; therapy; irritable bowel syndrome; diabetes; scleroderma; Parkinson's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A new method is provided for preparing a decapeptide of formula pGlu-His-Trp-Ser-Tyr-Xaa-Leu-Arg-Pro-Gly-NH2, in which a protected nonapeptide corresponding to the N-terminal of the peptide is first prepared on a benzhydrylamine resin, the Pro residue being attached to the resin via a photosensitive linker. The nonapeptide is cleaved from the resin by photolysis, the C-terminal is activated, and the product is coupled with glycinamide to add the Gly-NH2. The decapeptide is then deprotected. In the decapeptide, Xaa is Gly (giving gonadorelin; the present sequence), D-2-Nal or D-Trp.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Prepn. of luteinising hormone and follicle stimulating hormone releasing peptide(s) - by cleaving a protected nona-peptide resin by photolysis to remove the support, coupling with glycinamide and
                                      luteinising hormone releasing hormone; follicle stimulating; FSH; gonadorelin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 10;
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100.0%; Pred. No. 0.00032;
tive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                     (BOEH ) BIO-MEGA/BOEHRINGER INGELHEIM RES INC.
                                                                                                                                                                                    'note= "pyroglutamic acid"
                                                                                                                                           Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAR86845 standard; peptide; 10 AA
                                                                                                                                                                                                                             /note= "Gly-NH2"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 1; Page ?; 18pp; English.
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                                                                                                                                                                                                                                                                                                                                                   88CA-0566195.
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(first entry)
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les 10; Conservative
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                                                                                                                                                Key
Modified-site
                                                                                                                                                                                                        Modified-site
    LHRH peptide
                                                                                                                                                                                                                                                                                                                                                      06-MAY-1988;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gauthier JA;
                                                                                                                                                                                                                                                                      CA1335403-C
                                                                                                                                                                                                                                                                                                             25-APR-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        25-MAR-2003
22-MAR-1996
                                                                                                       Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic immunogenic peptides are provided in which a universal immune stimulator is linked to a peptide or protein hapten containing B cell and/or cytotoxic T lymphocyte epitopes, giving a product which causes potent immune responses to the coupled peptide or protein. The stimulator consists of (A) a promiscuous helper T cell epitope (Th) which elicits an immune response to the coupled peptide in members of a heterogeneous population expressing diverse HLA phenotypes, and (B) an adjuvant peptide sequence from the invasin protein of Yersinia. Spacer amino acid sequence from the invasin protein of Yersinia. Invasin and Th domains and between the immune stimulator and hapten components. When the hapten is LHRH, then optionally the invasin domain can be omitted from the immune stimulator component.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present sequence represents an LHRH hapten which can be attached to the stimulator to provide a potent vaccine for treating e.g. prostatic hyperplasia, androgen-dependent carcinoma, prostatic carcinoma, testicular carcinoma, endometriosis, benign uterine tumours, recurrent functional ovarian cysts, (severe) premenstrual syndrome or oestrogen-dependent breast cancer, or for
                                                                                                     Helper T cell epitope; universal immune stimulator; invasin; har
vaccine; LHRH; luteinising hormone releasing hormone; prostate;
androgen-dependent carcinoma; antitumour; infertility.
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0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Immunogenic luteinising hormone releasing hormone peptide(s) that suppress LHRH activity in males and females
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                                                                LHRH hapten for attachment to universal immune stimulator.
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                                                                                                                                                                                                                                                                                                                                                      93US-0057166
94US-0229275
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(updated)
(first entry)
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EHWSYGLRPG 10
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Ladd AE, Wang CY,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                10 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                (LADD/) LADD A E (WANG/) WANG C Y (ZAMB/) ZAMB T.
                                                                                                                                                                                                                                                                                                                                                      27-APR-1993;
14-APR-1994;
                                                                                                                                                                                                                                                                                                               28-APR-1994;
                                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                                                             WO9425060-A1
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    25-MAR-2003
                                                                                                                                                                                                                                                                      10-NOV-1994.
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                     10-SEP-1995
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Gaps

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Matches

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New gonadotropin releasing hormone preprohormone DNA - used to develop prods. for regulation of reproductive function and diagnosis of reproductive capacity and disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Bradykinin; N-benzylglycine; agonist, receptor study, antagonist, achiral; analgesic; luteinising hormone-releasing hormone; LHRH;
                                                                                                                                                                                                                                                                                                                                                                                                                               8 Different forms of GnRH (given in AAR75152-59) have previously
                                                                                                                                                                                                                                                                                                                                                                                                                                                       been isolated from vertebrate species. A precursor for an additional form of GnRH, (Ser8)-GnRH (AAR75151), has now been
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            100.0%; Score 63; DB 16;
100.0%; Pred. No. 0.00032;
Live 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Luteinising hormone-releasing hormone (LH-RH).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /note= "C-terminal amide"
                                                                                                                                                                             (STRD ) UNIV LELAND STANFORD JUNIOR. (UYOR-) UNIV OREGON HEALTH SCI. (UYOR-) UNIV OREGON STATE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAW65201 standard; peptide; 10 AA.
                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Fig.1a; 85pp; English
                                                                                                          94WO-US12763.
                                                                                                                                            93US-0147771.
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94US-0335202
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   10; Conservative
                                                                                                                                                                                                                                                    Adelman JP, Fernald RD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (REGC ) UNIV CALIFORNIA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            10 AA;
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Modified-site
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                                                                                                                                            05-NOV-1993;
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07-NOV-1994;
                                   WO9512309-A1
                                                                                                        04-NOV-1994;
                                                                       11-MAY-1995.
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   Mammalia.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 15
AAW65201
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      This sequence represents naturally occuring gonadotropin releasing hormone (GRRH). Analogues of GRRH are represented by ARR66846-56.

Motility disorders, including functional bowel disease, can be treated by the administration of one of the GRRH analogues shown here (e.g. by the administration of one of the GRRH analogues shown here (e.g. production of reproductive hormones such as luteinisting hormone, progesterone and relaxin. Motility disorders are caused from abnormalities of the autonomic nervous system. The GRRH analogues are administered by injection (which may be intavenous, subcutaneous or intramuscular), or by a drug delivery system. The drug delivery system can comprise a drug implant with timed release, a masal spray or an injection of a long-lasting depo form. This method is used to alleviate symptoms such as nausea, vomiting, abdominal pain and caltered bowel habites. The sequences can be used to ctreat motility disorders in a wide variety of cher diseases including irritable bowel of syndroms, disperse, soleroderma and parkinson's disease.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ö
                                                                                                                                                                                                                                                                                                                                                                                                                                 Treating motility disorders associated with systemic lupus erythematosus - by admin. of gonadotropin releasing hormone analogue, to control nausea, vomiting, abdominal pain etc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            100.0%; Score 63; DB 16; Length 10; 100.0%; Pred. No. 0.00032;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Mismatches
                                                      /note= "pyroglutamic acid"
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; Column 3; 14pp; English
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                                                                                                          /note= "amidated"
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                                 label= OTHER
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90US-0626402.
91US-0744977.
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EHWSYGLRPG 10
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Best Local Similarity
Matches 10; Conserv
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Key
Modified-site
                                                                       Cleavage-site
Modified-site
                                                                                                                                                                                                                 19-OCT-1992;
                                                                                                                                                                                                                                                   19-OCT-1992;
14-DEC-1990;
                                                                                                                                                                                                                                                                                      14-AUG-1991;
                                                                                                                                                                             18-JUL-1995
                                                                                                                                                                                                                                                                                                                                                             Mathias JR;
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Gaps

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Length 10; 0; Indels

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The invention relates to the obtaining of a potent agonist or antagonist peptide by the replacement of selected amino acids with synthetic achiral amino acids. The present sequence represents a luteinising hormone-releasing hormone (LHRH).
                                                                             New bradykinin analogues contg. N-benzyl-glycine - useful as bradykinin agonists or antagonists, useful e.g. as analgesics
                                                                                                                                Disclosure; Columns 11-12; 15pp; English.
             Mitchell AR, Young JD;
                                          WPI; 1996-299898/30.
                                                                                                                                                                                                                                                 Sequence 10 AA;
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Search completed: November 17, 2003, 18:21:54 Job time : 41 secs 1 EHWSYGLRPG 10 |||||||||| 1 EHWSYGLRPG 10 g

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Query Match 100.0%; Score 63; DB 17; Length 10; Best Local Similarity 100.0%; Pred. No. 0.00032; Matches 10; Conservative 0; Mismatches 0; Indels

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Best Local Similarity
Matches 10; Conser
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US-10-117-364-1
                          TYPE: PRT
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Sequence 1, Appli
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Sequence 4, Appli
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Sequence 1, Appl
                                                                                                                              November 17, 2003, 18:23:09; Search time 29 Seconds (without alignments) 62.952 Million cell updates/sec
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Compugen Ltd.
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US-10-117-364-1

US-10-111-688-4

US-10-118-553-1

US-10-115-553-1

US-09-964-201A-32

US-09-964-201A-32

US-09-351-641-1349

US-10-351-641-1349

US-10-351-641-1349

US-10-551-641-1349

US-09-305-924-9

US-10-351-641-1349

US-10-351-641-1349

US-10-351-641-1349

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GenCore version (c) 1993 - 2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Published Applications AA:*
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Maximum Match 100%
Listing first 45 summaries
                                                                                          - protein search, using sw model
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Gapop 10.0 , Gapext 0.5
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Match Length DB
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63
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Perfect score:
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US-10-109-331-2
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US-09-746-945-1
US-09-810-601-8
US-09-810-601-8
US-09-810-601-10
US-09-810-601-10
US-09-810-601-11
US-09-810-601-12
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US-09-810-601-12
US-09-810-601-20
US-09-810-601-20
US-09-810-601-21
US-09-810-601-22
US-09-810-601-23
US-09-810-601-24
US-09-810-601-25
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ALIGNMENTS

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Sequence 28, Application US/09964201A

Sequence 28, Application US/09964201A

Publication No. US20030091575A1

GENERAL INFORMATION;

APPLICANT: Kenten, John H

APPLICANT: Pilon, Aprile L

APPLICANT: Pilon, Aprile L

APPLICANT: Polons, Gerald L

APPLICANT: Soberte Application No. US20030091575A1 09\026,276

CURRENT APPLICATION NUMBER: US/09/964,201A

CURRENT APPLICATION NOS: 35

SOFTWARE: PatentIn Ver. 2.0

SEQ ID NOS: 35

SEQ ID NOS: 35
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100.0%; Pred. No. 0.00075;
iive 0; Mismatches 0;
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; Sequence 1, Application US/10117364 ; Publication No. US20030181385A1 ; GENERAL INFORMATION: ; APPLICANT: Roeske, Roger W. TITLE OF INVENTION: LHRH Antagonist Peptides

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Gaps
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Publication No. US20030040482A1
GENERAL INFORMATION:
APPLICANT: Roseke, Roger W.
TITLE OF INVENTION: LHRH Antagonist Peptides
FILE REFERENCE: PPI-007CPUS
CURRENT APPLICATION NUMBER: US/10/115,553
CURRENT FILING DATE: 2002-04-02
PRIOR PILING DATE: EARLIER PILING DATE: 1998-04-06
PRIOR PILING DATE: EARLIER FILING DATE: 1998-04-06
PRIOR FILING DATE: EARLIER FILING DATE: 1998-06-07
NUMBER: OS SEQ ID NOS: 1
SOFTWARE: PETENTING DATE: 1995-06-07
SEQ ID NO 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     100.0%; Score 63; DB 15; Length 10; 100.0%; Pred. No. 0.00075;
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                                                                                                                                                                                                                                                          NAME/KEY: misc_feature
LOCATION: (1)._(1)
OTHER INFORMATION: Glu is modified with a pyro group.
FEATURE:
                                                                                                                                                                                                                                                                                                                                                             NAME/KEY: misc_feature; LOCATION: (10). (10) ... OTHER INFORMATION: Gly is modified with -NH2 group. US-10-184-126-1
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Fublication No. US2030044936A1
GENERAL INFORMATION
APPLICANT: Hwang, Jaulang
APPLICANT: Hsu, Chia-Tse
APPLICANT: Ting, Chun-Jen
TITLE OF INVENTION: PEPTIDE REPEAT INMUNOGENS
FILE REFERENCE: 08919-071001
CURRENT PAPLICATION NUMBER: US/10/122,483
CURRENT FILING DATE: 2002-07-15
PRIOR APPLICATION NUMBER: 09/412,558
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
PRIOR APPLICATION NUMBER: PCT/US99/09160
PRIOR FILING DATE: 1999-04-28
NUMBER OF SEQ 1D NOS: 1
SEQ ID NO 1
LENGTH: 10
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                                                                                                                                              TYPE: PRT ORGANISM: Artificial Sequence
                                                                                                                                                                                   FEATURE:
OTHER INFORMATION: synthetic
FEATURE:
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ORGANISM: Homo sapiens
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Matches 10; Conserv
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US-10-115-553-1
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US-10-122-483-1
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                        CURRENT APPLICATION NUMBER: US/10/117,364
CURRENT FILING DATE: 2002-04-05
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: US/08/973,378
PRIOR FILING DATE: EARLIER FILING DATE: 1998-04-06
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 08/480,494
PRIOR FILING DATE: EARLIER FILING DATE: 1995-06-07
NUMBER OF SEQ ID NOS: 1
SOFTWARE: PALENTIN Ver. 2.0
SEQ ID NO 1
LENGTH: 10
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TITLE OF INVENTION: PROPANE-1,3-DIONE DERIVATIVE
FILLE REPERENCE: 073475
CURRENT APPLICATION NUMBER: US/10/311,688
CURRENT FILING DATE: 2002-12-19
PRIOR PILING DATE: 2000-07-05
PRIOR PILING DATE: 2001-07-05
PRIOR FILING DATE: 2001-05-23
PRIOR FILING DATE: 2001-05-23
PRIOR FILING DATE: 2001-05-23
PRIOR PILING DATE: 2001-05-23
NUMBER OF SEQ ID NOS: 4
SOFTWARE: PATCHING DATE: 201-05-23
NUMBER OF SEQ ID NOS: 4
SOFTWARE: PATCHING DATE: 201-05-23
NUMBER OF SEQ ID NOS: 4
SOFTWARE: PATCHING DATE: 201-05-23
NUMBER OF SEQ ID NOS: 4
LENGTH: 10
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APPLICANT: ERACEY, Michael D.
TITLE OF INVENTION: PEG-LERH ANALOG CONJUGATES
FILE REFERENCE: EL-TAYAR=2A
CURRENT APPLICATION NUMBER: US/10/184,126
CURRENT FILING DATE: 2000-06-28
PRIOR APPLICATION NUMBER: US/09/698,134
FRICK FILING DATE: 2000-10-30
PRIOR FILING DATE: 1998-04-28
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Publication No. US20020183257A1
GENERAL INFORMATION:
APPLICANT: EL TAYAR, Nabil
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; Sequence 4, Application US/10311688; Publication No. US20030191164A1; GENERAL INFORMATION:
       FILE REFERENCE: PPI-007CPUS
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Best Local Similarity 100.
Matches 10; Conservative
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ORGANISM: Homo sapiens
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ORGANISM: Homo sapiens
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US-10-184-126-1
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Pred. No. 0.0023;
1; Mismatches 0; Indels
                                                                                                                                                                                                                                                     Indels
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APPLICANT: HARLAND, RICHARD
APPLICANT: HARLAND, RICHARD
APPLICANT: MANNS, JOHN G.
APPLICANT: MANNS, JOHN G.
APPLICANT: MANNS, JOHN G.
TITLE OF INVENTION: IMMUNIZATION AGAINST ENDOGENOUS
TITLE OF INVENTION: IMMUNIZATION AGAINST ENDOGENOUS
TITLE OF INVENTION: MOLECULES
NUMBER OF SEQUENCES: 6
CORRESPONDENCE ADDRESS:
ADDRESSER: ROBINS & ASSOCIATES
STREET: 90 MIDDLEFIELD ROAD, SUITE 200
CITY: MENLO PARK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CITY: MENLO PARK
STATE: CA
COUNTRY: USA
ZIP: 94025
COMPUTER REDABLE FORM:
MEDIUM TYPE: Floppy disk
COMPOTER: IBM PC compatible
CURRENT MADER: US/09/019,010
FILING DATE: 05-FEB-1998
CLASSIFICATION: 424
PRIOR APPLICATION: A24
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/036,883
FILING DATE: 05-FEB-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                      100.0%; Score 63; DB 15; 100.0%; Pred. No. 0.00075;
                                                                                                                                                                                                                                                   0; Mismatches
                     NUMBER OF SEQ ID NOS: 13
SOFTWARE: FastSEQ for Windows Version 4.0
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REGISTRATION NUMBER: 38,548

REFERENCE/DOCKET NUMBER: 9001-
TELEPOMUNICATION INFORMATION:
TELEPHONE: (550) 325-7812
INFORMATION FOR SEO ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 10 amino acids
TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 2, Application US/09019010
Patent No. US20010014330A1
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PRIOR FILING DATE: 1999-10-05
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Best Local Similarity 90.0
Matches 9; Conservative
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Best Local Similarity 100.
Matches 10; Conservative
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1 QHWSYGLRPG 10
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                                                                                                           TYPE: PRT
ORGANISM: Homo sapiens
US-10-122-483-1
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                                                                                        LENGTH: 10
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APPLICANT: Renten, John H
APPLICANT: Tramontano, Alfonso
APPLICANT: Tramontano, Alfonso
APPLICANT: Tramontano, Aprile L
APPLICANT: Lohnas, Gerald L
APPLICANT: Roberts, Steven F
TITLE OF INVENTION: HEAT-SHOCK FUSION-BASED VACCINE SYSTEM
FILE REPERENCE: U.S. Patent Application No. US20030091575A1 09\026,276
CURRENT APPLICATION NUMBER: US/09/964,201A
CURRENT FILING DATE: 2002-05-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 9, Application US/09305924A

| Sequence 9, Application US/09305924A
| Publication No. US20030091579A1
| COBREAL INFORMATION:
| APPLICANT: Stephen D. Acres
| TILE OF INVENTION: METHODS OF RAISING ANIMALS FOR MEAT PRODUCTION
| FILE REFERENCE: 9001-0048
| CURRENT APPLICATION NUMBER: US/09/305,924A
| CURRENT FILING DATE: 1999-05-05
| BARLIER APPLICATION NUMBER: US 60/084,217
| BARLIER FILING DATE: 1998-05-05
| NUMBER OF SEQ ID NOS: 14
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US-10-351-641-1143

Sequence 1143, Application US/10351641

Sequence 1143, Application O. US20030186874A1

GENERAL INFORMATION:

APPLICANT: Barney, S.

APPLICANT: Mouthis, G.

APPLICANT: Anwer, M.

APPLICANT: Anwer, M.

APPLICANT: Anwer, M.

TITLE OF INVENTION: PROPERTIES

TITLE OF INVENTION: PROPERTIES
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Pred. No. 0.0023;
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Sequence 32, Application US/09964201A Publication No. US20030091575A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                          NUMBER OF SEQ ID NOS: 35
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 32
LENGTH: 10
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Best Local Similarity 90.0°
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Best Local Similarity
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ORGANISM: Porcine
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US-09-305-924-9
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US-09-305-924-9
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TYPE: PRT
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RESULT 8

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APPLICANT: Anwer, M.
APPLICANT: Lambert, D.
TITLE OF INVENTION: HYBRID POLYPEPTIDES WITH ENHANCED PHARMACOKINETIC
TITLE OF INVENTION: HYBRID POLYPEPTIDES WITH ENHANCED PHARMACOKINETIC
TITLE OF INVENTION: HYBRID POLYPEPTIDES WITH ENHANCED PHARMACOKINETIC
TITLE REPRESENCE: 7822-100
CURRENT APPLICATION NUMBER: US/10/351,641
PRIOR PELLING DATE: 1999-07-09
PRIOR FILING DATE: 1999-07-09
PRIOR FILING DATE: 1999-07-09
PRIOR FILING DATE: 1999-05-20
PRIOR FILING DATE: 1999-05-20
PRIOR FILING DATE: 1998-05-20
NUMBER OF SEQ ID NOS: 1757
SEQ ID NO 1344
TENERAL PARESEQ FOR WINDOWS VERSION 3.0
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Sequence 1, Application US/09810601
Sequence 1, Application Stephen
TITLE OF INVENTION: Composotions and Methods for Treating Gonadotrophin
TITLE OF INVENTION: Related Illnesses
CURRENT PELLIGATION NUMBER: US/09/810,601
CURRENT FILING DATE: 2001-03-15
NUMBER OF SEQ ID NOS: 44
SOFTWARE: Patentin Ver. 2.1
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Pred. No. 0.0023;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity 90.0%;
Matches 9; Conservative
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Matches 9; Conservative
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       Guthrie, K.
Merutka, G.
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ORGANISM: human
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US-09-305-924-1
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LENGTH: 10
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APPLICANT: Barney, S.
APPLICANT: Barney, S.
APPLICANT: Guthrie, K.
APPLICANT: Guthrie, M.
APPLICANT: Lambert, M.
APPLICANTON: PROPERTIES
FILE REFERENCE: 7872-100
CURRENT PAPLICATION NUMBER: 0510-24
PRIOR FILING DATE: 1999-05-09
PRIOR FILING DATE: 1999-05-09
PRIOR PAPLICATION NUMBER: 09/315,304
PRIOR PAPLICATION NUMBER: 09/315,304
PRIOR PAPLICATION NUMBER: 09/315,304
PRIOR PAPLICATION NUMBER: 09/315,304
PRIOR PELING DATE: 1999-05-20
PRIOR PELING DATE: 1999-05-20
PRIOR PILING DATE: 1999-05-20
SOFTWARE: FastSEQ for Windows Version 3.0
SOFTWARE: FastSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
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Pred. No. 0.0023;
1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               95.2%; Score 60; DB 12; Length 10; 90.0%; Pred. No. 0.0023;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1; Mismatches
CURRENT FILING DATE: 2003-01-24
CURRENT FILING DATE: 2003-01-24
PRIOR APPLICATION NUMBER: 09/350,641
PRIOR FILING DATE: 1999-07-09
PRIOR FILING DATE: 1999-07-09
PRIOR FILING DATE: 1999-05-20
PRIOR FILING DATE: 1999-05-20
PRIOR FILING DATE: 1998-05-20
PRIOR FILING DATE: 1998-05-20
SOFTWARE: FastsEQ for Windows Version 3.0
SEQ ID NO: 1143
LENGTH: 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 11
2-10-31-641-1309
; Sequence 1309, Application US/10351641
; Publication No. US20030186874A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; Sequence 1344, Application US/10351641; Publication No. US20030186874A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                           FEATURE:

COTHER INFORMATION: Core polypeptide
US-10-351-641-1143
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         , OTHER INFORMATION: Core polypeptide US-10-351-641-1309
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                          ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity 90.0%;
Matches 9; Conservative
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Matches 9; Conservative
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US-10-351-641-1344
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; OTHER INFORMATION: Labeled Tyr with 1251 and substituted with D form of Trp US-10-311-688-3
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                                                                                                         TITLE OF INVENTION: METHODS OF RAISING ANIMALS FOR MEAT PRODUCTION FILE REFERENCE: 9001-0048
CURRENT APPLICATION NUMBER: US/09/305,924A
CURRENT FILING DATE: 1999-05-05
EARLIER FILING DATE: 1999-05-05
EARLIER FILING DATE: 1998-05-05
SOFTWARE: PRODUCTION NUMBER: US 60/084,217
SOFTWARE: PATENTING OFF: 2.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 92.1%; Score 58; DB 11; Length 10; Best Local Similarity 100.0%; Pred. No. 0.0048; Matches 9; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 3, Application US/10311688
Publication No. US20030191164A1
GENERAL INFORMATION:
GENERAL INFORMATION:
TITLE OF INVENTION: PROPANE.1,3-DIONE DERIVATIVE
FILE REFERENCE: 073475
CURRENT APPLICATION NUMBER: US/10/311,688
CURRENT FILING DATE: 2002-12-19
FRIOR APPLICATION NUMBER: UPA P. 2000-204425
FRIOR APPLICATION NUMBER: UPA P. 2001-153372
FRIOR FILING DATE: 2000-07-05
FRIOR FILING DATE: 2001-05-23
FRIOR FLING DATE: 2001-05-23
FRIOR FRIOR DATE: 2001-05-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FEATURE:
NAME/KEY: MOD_RES
LOCATION: (1)
TOTHER INFORMATION: Xaa is pyroglutamic acid
US-09-305-924-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: PRT
ORGANISM: Artificial Sequence
PEATURE:
                                          Stephen D. Acres
Richard Harland
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 87.3
Best Local Similarity 90.0
Matches 9; Conservative
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APPLICANT: Jack G. Manns
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ORGANISM: GnRH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-10-311-688-3
                                                                                                                                                                                                                                                                                                                                                                                                                    SEQ ID NO 1
LENGTH: 10
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Search completed: November 17, 2003, 18:28:17 Job time : 29 secs

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GenCore version 5.1.6 Copyright (c) 1993 - 2003 Compugen Ltd.
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OM protein - protein search, using sw model

November 17, 2003, 18:27:39 ; Search time 13.5 Seconds (without alignments) 64.112 Million cell updates/sec Run on:

US-09-462-089-2 58 1 HWSYGLRPG 9 Title: Perfect score: Sequence:

Scoring table:

283308 segs, 96168682 residues BLOSUM62 Gapop 10.0 , Gapext 0.5 Searched:

Total number of hits satisfying chosen parameters:

789

Minimum DB seq length: 0 Maximum DB seq length: 9

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database :

PIR 76:*
1: pir1:*
2: pir2:*
3: pir3:*
4: pir4:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

		di			SUMMARIES	
Result No.	Score	Query Match	, Length	03	ΙD	Description
	21	36.2	! ! !	2	839437	D-amino-acid oxida
7	19	32.8		7	A60139	fatty-acid synthas
٣	19	32.8	80	7	D47393	υ.
4	18	31.0		~	PT0268	Ig heavy chain CRD
2	18	31.0		7	PT0299	6
9	17	29.3		~	PH1618	Ig H chain V-D-J r
7	17	29.3		~	807205	ij
80		29.3	6.	N	S07204	litorin I - Austra
σ	17	29.3		~	D58503	translation elonga
10	17	29.3		7	PT0238	Ig heavy chain CRD
11	17	. 29.3		7	PH1591	Ig H chain V-D-J r
12	17	29.3	6	~	G41946	T-cell receptor ga
13	16	7.		N	PT0281	Ig heavy chain CRD
14	16	~		4	155382	hypothetical pepti
15	16	27.6		~	S07241	litorin - Rohde's
16	15	ഹ		~	A33098	244K exoantigen -
17	15	25.9		7	QDRB	delta sleep-induci
18	15	S		~	A11497	transaldolase (EC
19	15	ß		~	836850	Ig heavy chain V r
20	15	S		~	I49406	bone gla protein -
21	14	4		~	JN0862	peptidyl-dipeptida
22	14	24.1		4	179564	hypothetical TCL3
23	14	4		~	PT0581	T-cell receptor be
24	14			~	S19288	acylase - Kluyvera
25	14		8	~	S16324	hypothetical prote
26	14	24.1		~	S11078	glucose-6-phosphat
27	14		8	~	JS0318	leucokinin VIII -
28	14	24.1	80	N	PT0311	Ig heavy chain CRD
29	14	24.1	<u>۔</u> ص	7	PT0288	Ig heavy chain CRD

2 WSYG 5

alpha-2-macroglobu RPCH-related neuro Ig heavy chain CRD	ubiquitin - rat photosystem I 10.4 Leu-enkephalin - b	X 0)	1-Cell receptor be RNA-directed DNA p halo-toxin - Pseud T-cell recentor be	hypothetical colla Met-enkephalin-Arg platelet aggregati
			*	
S66636 A34626 PT0240	S47552 PQ0689 B61445	A61445 S53595 PT0572	PIU/14 A35890 A61049 PT0715	A35039 A60224 A44428
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0.44	4 10 10 1		, , , , ,	7 7 6 0
24 . 1 22 . 4 22 . 4	2222	222 222 4. 4. 4.	2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2	22.4 22.4 22.4
14 13	13 13	E E E E	1 1 1 1	113
30 31 32	8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8	9 M M M	2 4 4 4 2 0 4 4	. 4. 4. 4. 16. 4. 73

ALIGNMENTS

	RESULT 1
	Description of the control oxidase (EC 1.4.3.3) - Trigonopsis variabilis (fragment)
 -	C.)Species: irigonopars variantins C.)Date: 19-Mar-1997 #sequence_revision 05-Dec-1997 #text_change 07-May-1999
	C.Accession: 55943/ R;Schraeder, T., Andreesen, J.R. Eur. J. Biochem 218 735-744 1993
	A;Title: Evidence for the functional importance of Cys298 in D-amino acid oxidase from 1 A;Reference number: S39437; MUID:94094869; PMID:7903639
	A;Accession: S39437 A;Molecule type: protein
	A, Residues: 1-9 <sch> A, Experimental source: CBS 4095 C: Function:</sch>
	A;Description: oxidoreductase; catalyzes the oxidation of D-amino acids to their correst A;Note: reoxidation of the enzyme by molecular oxygen is accompanied by the release of T;C;Keywords: FAD; oxidoreductase
-	Query Match Best Local Similarity 80.0%; Pred. No. 2.8e+05; Matches 4; Conservative 0; Mismatches 1; Indels 0; Gaps 0;
	Qy S GLRPG 9
	Db 3 GHRPG 7
-	RESULT 2 A60139
	<pre>fatty-acid synthase (EC 2.3.1.85) - rabbit (fragment) C;Species: Oryctolagus cuniculus (domestic rabbit)</pre>
	C;Date: 22-Jan-1993 #sequence_revision 22-Jan-1993 #text_change 26-May-2000 C;Accession: A60139
	K;Hardie, D.G.; Dewart, K.B.; Altken, A.; McCartny, A.D. Biochim. Biochiw. Acta 828, 380-382, 1985
	A;Title: Amino acid sequence around the reactive serine residue of the unloesterase dom A;Reference number: A60139; MUID:85175165; PMID:3921056 A:Arression: A60139
	A;Moccule type: protein A:Moclecule type: protein A:Residues: 1-7 <har></har>
	C;Superfamily: rat fatty-acid synthase; 3-oxoacyl-[acyl-carrier-protein] synthase I homydrolase homology; [acyl-carrier-protein] Sydrolase homology; [acyl-carrier-protein] S
	C;Keywords: acyltransferase; carrier protein; coenzyme A; homodimer; multifunctional en· F;5/Active site: Ser (of oleoyl-[acyl-carrier-protein] hydrolase) #status experimental
	Query Match 32.8%; Score 19; DB 2; Length 7;

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PHIG18
19 H chain V-D-J region (clone B-less 33) - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Species: Mus musculus (house mouse)
C;Date: 02-Jun-1994 #sequence_revision 02-Jun-1994 #text_change 17-Mar-1999
C;Date: 02-Jun-1994 #sequence_revision 02-Jun-1994 #text_change 17-Mar-1999
C;Accession: PHIS18
B;Levinson, D.A.; Campos Torres, J.; Leder, P.
J. Exp. Med. 178, 317-329, 1993
J. Exp. Med. 178, 317-329, 1993
A;Title: Molecular characterization of transgene-induced immunodeficiency in B-less mice
A;Reference number: PHIS80; MUID:93301609; PMID:8315387
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RiAnastasi, A.; Montecucchi, P.; Angelucci, F.; Erspamer, V.; Endean, R. Experientia 33, 1289, 1977

Experientia 33, 1289, 1977

Experientia 34, 1289, 1977

Experientia 35, 1289, 1977

Experientia 36, 1289, 1977

A; Fitle: Glu(OMb) (2)-litorin, the second bombesin-like peptide occurring in methanol ext A; Reference number: S07205; MUID: 78003546; PMID: 908397

A; Accession: S07205
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Experientia 31, 510-511, 1975
A;Title: Aminoacid composition and sequence of litorin, a bombesin-like nonapeptide from
A;Reference number: S07204; MUID:75187011; PMID:1140241
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C;Date: 12-Feb-1993 #sequence_revision 12-Mar-1993 #text_change 18-Aug-2000
C;Accession: S07205
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C;Date: 12-Feb-1993 #sequence_revision 12-Mar-1993 #text_change 18-Aug-2000
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A,Residues: 1-9 <ANA>
(Superfamily: gastrin-releasing peptide
C,Keywords: amidated carboxyl end; neuropeptide; pyroglutamic acid
F;1/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental
F;9/Modified site: amidated carboxyl end (Met) #status experimental
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A,Molecule type: protein
C,Superfamily: gastrin-releasing peptide
C,Superfamily: gastrin-releasing neuropeptide; pyroglutamic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 17; DB 2; Length 9;
Pred. No. 2.8e+05;
1; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       29.3%; Score 17; DB 2; L6 100.0%; Pred. No. 2.8e+05; Itive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     litorin 2-Glu - Australian tree frog (Litoria aurea)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A; Experimental source: bone marrow pre-B lymphocyte C; Keywords: immunoglobulin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  litorin I - Australian tree frog (Litoria aurea)
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Best Local Similarity 100.
Matches 3; Conservative
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Best Local Similarity 50.0
Matches 2; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A; Molecule type: DNA A; Residues: 1-8 < LEV >
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2 WSYG 5
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                                4
                                                                                       WDY 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             6 YGL 8
                                WSY
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                                                                                                                                                                                                            C. Jate: 16-Feb-1994 #sequence_revision 18-Nov-1994 #text_change 28-Apr-1995
C. Accession: D47393
C. Accession: D47393
S. Johnsen, A.H.; Scott, A.G.; Yu, C.G.; Yagi, K.J.; Tobe, S.S.; Thorpe, A. Proc. Natl. Acad. Sci. U.S.A. 90, 2456-2460, 1993
A. Title: Callatoscatins: neuropeptides from the blowfly Calliphora vomitoria with sequent A; Reference number: A47393; MUD:93211980; PMID:8460157
A; Accession: D47393
A; Status: proliminary
A; Molecule type: protein
A; Residues: 1-8 <a href="https://documentalsource: thoracic ganglia">https://documentalsource: thoracic ganglia</a>
A; Note: sequence extracted from NCBI backbone (NCBIP:128479)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         C; Accession: PT0268
R; Yamada, M.; Wasserman, R.; Reichard, B.A.; Shane, S.; Caton, A.J.; Rovera, G.
J. Exp. Med. 173, 395-407, 1991
A; Title: Preferential utilization of specific immunoglobulin heavy chain diversity and A; Reference number: PT0222; MUID:91108337; PMID:1899102
A; Reference number: PT0222; MUID:91108337; PMID:1899102
A; Recidine: 1-9 < YAM>
A; Residues: 1-9 < YAM>
A; Residues: 1-9 < YAM>
A; Residues: heterotetramer; immunoglobulin
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CiSpecies: Homo sapiens (man)
CiSpecies: Homo sapiens
CiAccession: PT0299
RiYamada, M.; Wasserman, R.; Reichard, B.A.; Shane, S.; Caton, A.J.; Rovera, G. J. Exp. Med. 173, 395-407, 1991
A;Title: Preferential utilization of specific immunoglobulin heavy chain diversity A;Reference number: PT0222; MuID:91108337; PMID:1899102
A;Reference number: PT0222; MuID:91108337; PMID:1899102
A;Reference number: DNA
A;Reference number: Inmunoglobulin
C;Keywords: heterotetramer; immunoglobulin
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                                                                                                                                                 neuropeptide callatostatin 4 - bluebottle fly (Calliphora vomitoria)
C;Species: Calliphora vomitoria
C;Date: 16-Fb-1994 #sequence_revision 18-Nov-1994 #text_change 28-Apr-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        C;Species: Homo sapiens (man)
C;Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change 16-Aug-1996
C;Accession: PT0268
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             32.8%; Score 19; DB 2; Length 8; 60.0%; Pred. No. 2.8e+05; ive 2; Mismatches 0; Indels
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Matches 3; Conservative 0; Mismatches
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Best Local Similarity
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4 YSFGL 8
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Gaps .; 0

Length 9;

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C,Accession: G41946
Whetsell, M.; Mosley, R.L.; Whetsell, L.; Schaefer, F.V.; Miller, K.S.; Klein, J.R. Mol. Cell. Biol. 11, 5902-5909, 1991
A,Title: Rearrangement and junctional-site sequence analyses of T-cell receptor gamma ge A,Reference number: A41946, MUID:92049316; PMID:1658619
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  R;Yamada, M.; Wasserman, R.; Reichard, B.A.; Shane, S.; Caton, A.J.; Rovera, G. J. Exp. Med. 173, 395-407, 1991
A;Title: Preferential utilization of specific immunoglobulin heavy chain diversity and A;Reference number: PT0222; MUID:91108337; PMID:1899102
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C;Species: Homo sapiens (man)
C;Date: 16-Apr-1999 #sequence_revision 16-Apr-1999 #text_change 20-Apr-2000
C;Accession: I55382
R;Dawson, S:J: Wiman, B:; Hamsten, A.; Green, F:; Humphries, S.; Henney, A.M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      C;Species: Mus musculus (house mouse)
C;Date: 03-Feb_1994 #sequence_revision 03-Feb-1994 #text_change 07-May-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ig heavy chain CRD3 region (clone 4-91C) - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change 16-Aug-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A, Status: preliminary, not compared with conceptual translation A, Molecule type: DNA
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   A;Reference number: PH1580; MUID:93301609; PMID:8315387
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           - mouse (fragment)
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Best Local Similarity 100.0%; Pred. No. 2.8e+05;
Matches 3; Conservative 0; Mismatches 0;
                                                                                                                                                                                                    29.3%; Score 17; DB 2; Le
100.0%; Pred. No. 2.8e+05;
tive 0; Mismatches 0;
                                                                                                                  bone marrow pre-B lymphocyte
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A;Residues: 1-5 <YAM>
A;Experimental source: B lymphocyte
C;Keywords: heterotetramer; immunoglobulin
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C;Keywords: T-cell receptor
                              A;Accession: PH1591
A;Molecule type: DNA
A;Residues: 1-9 < LEV>
A;Experimental source: bone
C;Keywords: immunoglobulin
                                                                                                                                                                                                                         Best Local Similarity
Matches 3; Conser
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                                                                                                                                                                                                                                                                                                                                                                                                                                   C;Species: unidentified bacterium
C;Species: unidentified bacterium
C;Species: O7-Peb-1997 #sequence_revision 07-Peb-1997 #text_change 28-May-1999
C;Accession: D5803
R;Binette, J.P.; Binette, M.B.
Sibminete, J.P.; Binette, M.B.
R;Binette, J.P.; Binette, M.B.
R;Binette, J.P.; Binette, M.B.
A;Becription: The protein Sequence Database, October 1996
A;Becription: The protein of kidney and gallbladder stones.
A;Recience number: A58501
A;Accession: D5803
A;Status: protein
A;Status: protein
A;Status: protein
A;Residues: 1-9 < EBIN>
A;Residues: 1-9 < EBIN>
A;Experimental source: human bile and stones
C;Superfamily: translation elongation factor Tu; translation elongation factor Tu homolc
C;Keywords: GTP binding
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C;Species: Mus musculus (house mouse)
C;Date: 02-Jun-1994 #sequence_revision 02-Jun-1994 #text_change 17-Mar-1999
C;Accession: PH1591
R;Levinson, D.A.; Campos-Torres, J.; Leder, P.
J. Exp. Med. 178, 317-329, 1993
A;Title: Molecular characterization of transgene-induced immunodeficiency in B-less mice
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         C;Accession: PT0238
R;Yamada, M; Wasserman, R.; Reichard, B.A.; Shane, S.; Caton, A.J.; Rovera, G.
J. Exp. Med. 173, 395-407, 1991
A;Title: Preferential utilization of specific immunoglobulin heavy chain diversity and A;Reference number: PT0222; MUID:91108337; PMID:1899102
A;Accession: PT0238
A;Accession: PT0238
A;Molecule type: DNA
A;Residues: 1-9 <YAM>A;Residues: 1-9 <YAM>A;Residues: los <YAM>A;Residues: heterotetramer; immunoglobulin
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C;Species: Homo sapiens (man)
C;Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change 16-Aug-1996
C;Accession: PT0238
                                                                                                                                                                                                                                                                                                                                                                                                        translation elongation factor EF-Tu - unidentified bacterium (fragment)
C;Species: unidentified bacterium
C;Date: 07-Feb-1997 #sequence_revision 07-Feb-1997 #text_change 28-May-1999
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F;1/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental F;9/Modified site: amidated carboxyl end (Met) #status experimental
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Best Local Similarity 50.03
Matches 2; Conservative
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J. Biol. Chem. 268, 10739-10745, 1993
A.Title: The two allele sequences of a common polymorphism in the promoter of the plasmi A.Reference number: I55382; MUID:93266509; PMID:8388372
A.Accession: I55382
A.Accession: I55382
A.Molecule type: DNA
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A.Molecule type: DNA
A.Molecule type: DNA
A.Coss.references: GB:M91557; NID:g190020; PIDN:AAA60110.1; PID:g190021
C.Comment: This is the hypothetical translation of a sequence from the PAII gene promote C.Comment: This is the hypothetical translation of A sequence from the PAII gene promote A.Geneic GBB:NA1
A.Coss-references: GBB:120297; OMIM:173360
A.Mop position: 7q21.3-7q22
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C.Species: Phyllomedias robde; (Robde's leaf frog)
C.Date: 12-Peb-1993 #sequence_revision 12-Mar-1993 #text_change 18-Aug-2000
C.Date: 12-Reb-1993 #sequence_revision 12-Mar-1993 #text_change 18-Aug-2000
C.Accession: 807241
A.Title: Robdei-litorin: a new peptide from the skin of Phyllomedusa robdei.
A.Accession: 807241
A.Accession: 80
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Copyright (c) 1993 - 2003 Compugen Ltd
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Eur. J. Biochem. 250:727-734(1997)
-!- FUNCTION: MAY ACT AS A NEUROTRANSMITTER OR NEUROMODULATOR.
-!- SIMILARITY: BELONGS TO THE ALLATOSTATIN FAMILY.
Neuropeptide; Amidation; Multigene family.
MOD RES
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AMIDATION (POTENTIAL).
SEQÜENCE 5 AA; 586 MW; 672879D5AB300000 CRC64;
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Eur. J. Biochem. 250:727-734 (1997).
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-!- SIMILARITY: BELONGS TO THE ALLATOSTATIN FAMILY.
Neuropeptide; Amidation; Multigene family.
MOD_RES 8 8 AMIDATION.
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Bukaryota; Metazoa, Arthropoda; Crustacea; Malacostraca;
Eumalacostraca; Eucarida; Decapoda; Plecoyemata; Brachyura;
Bubrachyura; Portunoidea; Portunidae; Carcinus.
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Pred. No. 1.3e+05;
2; Mismatches 0; Indels
                                                          8 AA; 813 MW; 7C286B45AB476878 CRC64;
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30-MAY-2000 (Rel. 39, Last sequence update)
30-MAY-2000 (Rel. 39, Last annotation update)
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30-MAY-2000 (Rel. 39, Last sequence update)
30-MAY-2000 (Rel. 39, Last annotation update)
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-!- FUNCTION: MAY ACT AS A NEUROTRANSMITTER OR NEUROMODULATOR.
-!- SIMILARITY: BELONGS TO THE ALLATOSTATIN FAMILY.
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allatostatin superfamily in the shore crab Carcinus maenas.";
Eur. J. Biochem. 250:727-734(1997).
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-i- SIMILARITY: BELONGS TO THE ALLATOSTATIN FAMILY.
Neuropeptide; Amidation; Multigene family.
MOD RES 8 8 AMIDATION.
SEQÜENCE 8 AA; 811 MW; 922879D5AB47687D CRC64;
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Carcinustatin 18.

Carcinus maenas (Common shore crab) (Green crab).

Bukaryota; Metazoa; Arthropoda; Crustacea; Malacostraca;

Bumalacostraca; Bucarida; Decapoda; Plecyemata; Brachyura;

Bubrachyura; Portunoidea; Portunidae; Carcinus.
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                                                                                                                                                                                                                                                    Length 8;
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MOD RES 8 AMIDATION (POTENTIAL)
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                                                                                                                                                                                                                                                    Score 19; DB 1;
Pred. No. 1.3e+05;
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30-MAY-2000 (Rel. 39, Last sequence update)
30-MAY-2000 (Rel. 39, Last annotation update)
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Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha; Oestroidea;
Calliphoridae; Calliphora.
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TISSUE=Larva;
MEDLINE=98054519; PubMed=9392829;
BUVE H., Johnson A.H., Maestro J.-L., Scott A.G., Winstanley D.,
Davey M., East P.D., Thorpe A.;
"Lepidopteran peptides of the allatostatin superfamily.";
Peptides 18:1301-1309(1997).
-: SIMILARITY: BELONGS TO THE ALLATOSTATIN FAMILY.
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"Isolation and identification of multiple neuropeptides of the
                                    allatostatin superfamily in the shore crab Carcinus maenas.";
Eur. J. Biochem. 250:727-734(1997).
-!- FUNCTION: MAY ACT AS A NEUROTRANSMITTER OR NEUROMODULATOR.
-!- SIMILARITY: BELONGS TO THE ALLATOSTATIN FAMILY.
Neuropeptide; Amidation; Multigene family.
MOD RES 8 8 AMIDATION (POTENTIAL).
SEQÜENCE 8 AA; 919 MW; C82879D5AB569AB5 CRC64;
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Eukaryota, Metazoa, Arthropoda, Hexapoda, Insecta, Pterygota, Neopera, Endoprery Sandoperygota, Lepidoptera, Glossata, Ditrysia, Tortricoidea, Tortricoidea, Tortricoidea, Cortricoidea, Cortriolea, Cort
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 32.8%; Score 19; DB 1; Length 8; Best Local Similarity 60.0%; Pred. No. 1.3e+05; Matches 3; Conservative 2; Mismatches 0; Indels
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30-MAY-2000 (Rel. 39, Last sequence update)
30-MAY-2000 (Rel. 39, Last annotation update)
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MEDLINE=93211980; PubMed=8460157;
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Best Local Similarity
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                                                                                                                       DOVE H., Thorpe A.,
"Distribution and functional significance of Leu-callatostatins in
the blowfly Calliphora vomitoria.",
Cell Tissue Res. 276:367-379(1994)
-!- FUNCTION: MAY ACT AS A NEUROTRANSMITTER OR NEUROMODULATOR AND PLAY
A ROLE IN THE INTERRATION OF INFORMATION WITHIN THE BRAIN MAY BE
INVOLVED IN THE CONTROL OF VISCERAL MUSCLES DUE TO ITS ABILITY TO
BEHAVE AS POTENT INHIBITORS OF PERISTALIC MOVEMENTS. MAY ALSO
FULFILL A NEUROHORMONAL ROLE ON MUSCLES OF THE GUT AND HEART.
-!- TISSUE SPECIFICITY: BRAIN, SUBOESOPHAGEAL GANGLION, RETROCEREBRAL
COMPLEX, THORACICO-ABDOMINAL GANGLION, PERIPHERAL NEUROSECRETORY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Duve H., Johnsen A.H., Maestro J.-L., Scott A.G., Winstanley D., Davey M., East P.D., Thorpe A.; "Lepidopteran peptides of the allatostatin superfamily."; Peptides 18:1301-1309(1997).

-: SIMILARITY: BELONGS TO THE ALLATOSTATIN FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Cydiastatin 4.
Cydia pomonella (Codling moth).
Cydia pomonella (Codling moth).
Bukaryota, Metazoa, Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Lepidoptera; Glossata; Ditrysia;
Tortricoidea; Tortricidae; Olethreutinae; Cydia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 19; DB 1; Length 8; Pred. No. 1.3e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 8;
with sequence homology to cockroach allatostatins.";
Proc. Natl. Acad. Sci. U.S.A. 90:2456-2460(1993).
                                                                                                                                                                                                                                                                                                                                                                                                                           SYSTEM AND INTESTINE.
-!- SIMILARITY: BELONGS TO THE ALLATOSTATIN FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               8 AMIDATION.
8 AA; 910 MW; 922879D5AB47740D CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       8 AA; 954 MW; D32879D5AB47740A CRC64;
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30-MAY-2000 (Rel. 39, Last sequence update)
30-MAY-2000 (Rel. 39, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              8 AA.
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OR N.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TISSUE=Larva;
MEDLINE=98054539; PubMed=9392829;
                                                                                                 MEDLINE=94291167; PubMed=8020069;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TKCI_CALVO
ID TKCI_CALVO STANDARD;
ID TKCI_CALVO STANDARD;
ID TKCI_CALVO STANDARD;
ID TNCV-1995 (Rel. 32, Created)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       32.8%;
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60.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    3; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     3; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Neuropeptide; Amidation.
MOD RES 8 8
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STANDARD;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NCBI_TaxID=9606;
                                                                                                                                                                                                                                        NCBI_TaxID=8371;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2 WSYG 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                         Litoria aurea.
                                             01-NOV-1988
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               HUMAN
LITAU
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SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 P30092;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           UPA6_HUMAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
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DDT THE SET TH
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                                                                                                                                                                                                                                                     raessel D.R.;

"Callitachykinin I and II, two novel myotropic peptides isolated from the blowfly, Calliphora vomitoria, that have resemblances to tachykinins.";

Peptides 15:61-768(1994).

-! FUNCTION: MYOACTIVE PEPTIDE.

-! SUBCELLULAR LOCATION: Secreted.

-! SUBCELLULAR LOCATION: Acceted.

-! SUBCELLULAR LOCATION: Acceted.
                                                                                            Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha; Oestroidea;
Calliphoridae; Calliphora.
                                                                                                                                                                                                                                        MEDLINE=95075727; PubMed=7984492;
Lundquist C.T., Clottens F.L., Holman G.M., Nichols R., Nachman R.J.,
Naessel D.R.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDLINE=90184489; PubMed=2311766; Schoofs L., Holman G.M., Hayes T.K., Nachman R.J., de Loof A.; Schoofs L., Holman G.M., Hayes T.K., Nachman R.J., de Loof A.; Schoofs L., Holman II, two novel insect neuropeptides with homology to peptides of the vertebrate tachykinin family."; FEBS Lett. 261:397-401(1990).

-! FUNCTION: MYOACTIVE PEPTIDE. STIMULATES THE CONTRACTION OF THE OVIDUCT AND FOREGUT.

-! SUBCELLULAR LOCATION: Secreted.
-! SIMILARITY: SOME SIMILARITY TO TACHYKININS.
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Eukaryota; Metazoa, Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Orthopteroidea; Orthoptera; Caelifera; Acridomorpha;
Acridoidea; Acrididae; Oedipodinae; Locusta.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      32.8%; Score 19; DB 1; Length 9; 75.0%; Pred. No. 1.3e+05; cive 1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  32.8%; Score 19; DB 1; Length 9; 75.0%; Pred. No. 1.3e+05;
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MOD RES 9 9 AMIDATION.
SEQUENCE 9 AA; 939 MW; 2389C86B59C865A7 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   9 AA; 981 MW; 2417C86B59CDC1B7 CRC64;
01-NOV-1995 (Rel. 32, Last sequence update)
15-SEP-2003 (Rel. 42, Last annotation update)
Callitachykinin I.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 01-APR-1990 (Rel. 14, Created)
01-APR-1990 (Rel. 14, Last sequence update)
15-SEP-2003 (Rel. 42, Last annotation update)
Locustatachykinin I (TK-I).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       9 AA.
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                                                                   Calliphora vomitoria (Blue blowfly).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3; Conservative
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                                                                                                                                                                                                                    AND SYNTHESIS.
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Matches 3; Conserv
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Best Local Similarity
Matches 3; Conserv
                                                                                                                                                                       NCBI_TaxID=27454;
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YGVR 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  4 YGLR 7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE.
TISSUE=Brain;
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RESULT 12 LITO_LITAU

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Anastasi A., Montecucchi P.C., Angelucci F., Erspamer V., Endean R.; "Glu(OMe)3-litorin, the second bombesin-like peptide occurring in methanol extracts of the skin of the Australian frog Litoria aurea."; Experientia 33:1289-1289(1977).
-! SUBCELLULAR LOCATION: Secreted.
-! TISSUE SPECIFICITY: Skin.
-! SIMILARITY: BELONGS TO THE BOMBESIN/NEUROMEDIN B/RANATENSIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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InterPro; IPR00874; Bombesin.
Pfam; PF02044; Bombesin; 1.
PROSITE; PS00287; BOMBESIN; 1.
Amphibian defense peptide; Bombesin family; Amidation; Methylation;
                                                                                           Litoria aurea (Green and golden bell frog).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Amphibia; Batrachia; Anura; Neobatrachia; Bufonoidea; Hylidae;
Pelodryadinae; Litoria.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                              TISSUE=Skin secretion;
MEDLINE=75187011; PubMed=1140241;
Anastasi A., Erspamer V., Endean R.;
"Aminoacid composition and sequence of litorin, a bombesin-like nonapeptide from the skin of the Australian leptodactylid frog
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01-APR-1993 (Rel. 25, Last sequence update)
16-OCT-2001 (Rel. 40, Last annoration update)
Unknown protein from 2D-page of plasma (Spot 14) (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PYRROLIDONE CARBOXYLIC ACID. METHYLATION (PARTIAL). AMIDATION. D7CCCIE862CDC366 CRC64;
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Pred. No. 1.3e+05;
1: Mismatches 1; Indels
                          01-FEB-1994 (Rel. 28, Last sequence update)
15-SEP-2003 (Rel. 42, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              9 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1; Mismatches
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MEDLINE=93092937; PubMed=1459097;
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MEDLINE=78003546; PubMed=908397;
(Rel. 09, Created)
(Rel. 28, Last seq
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE (METHYLATED VARIANT).
                                                                                                                                                                                                                                                                                                                                                                                                                                    Experientia 31:510-511(1975).
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MOD_RES 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           29.3%;
50.0%;
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Query Match
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-!- MISCELLANEOUS: ON THE 2D-GEL THE DETERMINED PI OF THIS UNKNOWN PROTEIN IS: 5, ITS MW IS: 48 kDa.
                                                                                                                              ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  "Isolation and identification of multiple neuropeptides of the allatostatin superfamily in the shore crab Carcinus maenas."; bur. J. Biochem. 250:77-734(1997).
-!- FUNCTION: MAY ACT AS A NEUROTRANSMITTER OR NEUROMODULATOR.
-!- SIMILARITY: BELONGS TO THE ALLATOSTATIN FAMILY.
Neuropeptide; Amidation; Multigene family.
MOD RES
7 AA; 770 MW; 672879CDESDDB70 CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                              TISSUE-Cerebral ganglion, and Thoracic ganglion;
MEDLINE=98121193; PubMed=9461295;
Duve H., Johnsen A.H., Maestro J.-L., Scott A.G., Jaros P.P.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE.
TISSUES-cerebral ganglion, and Thoracic ganglion;
MEDLINE-89121193; PubMed-9461295;
Duve H., Johnsen A.H., Maestro J.-L., Scott A.G., Jaros P.P.,
                                                                                                                                                                                                                                                                                                                                                                     Eumalacostraca; Eucarida; Decapoda; Plecyemata; Brachyura;
Eubrachyura; Portunoidea; Portunidae; Carcinus.
NCBI_TaxID=6759;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Carcinustatin 3.
Carcinus maenas (Common shore crab) (Green crab).
Bukaryota; Metazoa; Arthropoda; Crustacea; Malacostraca;
Eumalacostraca; Bucarida; Decapoda; Pleocyemata; Brachyura;
Bubrachyura; Portunoidea; Portunidae; Carcinus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   27.6%; Score 16; DB 1; Length 7; 40.0%; Pred. No. 1.3e+05; ative 3; Mismatches 0; Indels
                                                                                                                           1; Indels
                                                                                                                                                                                                                                                                                                                             Carcinustatin 2.
Carcinus maenas (Common shore crab) (Green crab).
Eukaryota; Metazoa; Arthropoda; Crustace; Malacostraca;
                                                                                               29.3%; Score 17; DB 1; Length 9; 75.0%; Pred. No. 1.3e+05; tive 0; Mismatches 1; Indels
                                                                    9 AA; 935 MW; 5282F2CAA8676447 CRC64;
                                                                                                                                                                                                                                                                                      30-MAY-2000 (Rel. 39, Created)
30-MAY-2000 (Rel. 39, Last sequence update)
30-MAY-2000 (Rel. 39, Last annotation update)
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                            SWISS-2DPAGE; P30092; HUMAN.
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les 2; Conservative
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3 YAFGL 7
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RI allatostatin superfamily in the shore crab Carcinus maenas.";

RL BLY. J. Biochem. 250:727-734(1997).

CC -!- FUNCTION: MAX AT AS A NEUROTRANSMITTER OR NEUROMODULATOR.

CC -!- SIMILARITY: BELONGS TO THE ALLATOSTATIN FAMILY.

KW Neuropeptide; Multigene family.

SQ SEQUENCE 7 AA; 796 MW; 672879CDC8476B70 CRC64;

Query Match

Best Local Similarity 40.0%; Pred. No. 1.3e+05;

Matches 2; Conservative 3; Mismatches 0; Indels 0; Gaps

QY 2 WSYGL 6

:::|| Db 3 YAFGL 7

Search completed: November 17, 2003, 18:30:10

Job time: 10 secs
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056140 syrenecocc
052062 bacillus me
015901 homo sapien
002032 lytechinus
091478 canis famil
09x871 caryctolagus
081802 zea may8 (m
P82079 limnodynast
                                                                                                                                                                                                                                                          999193 pseudomonas
031415 gallus gall
094hb1 microcebus
094hb2 microcebus
094xe6 tectocoris
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012096 caprine art
012102 caprine art
012102 caprine art
012098 caprine art
012098 caprine art
            Q62530 mus spretus
Q65711 berne virus
Q9prj4 lepisosteus
O85406 coxiella bu
                                                                          Q98tu5 xenopus lae
O49223 glycine max
                                                                                                         009258 synechococc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Vuorio E.;
"Evidence for insufficient chondrocytic differentiation during repair of full-thickness defects of articular cartilage.";
Matrix Biol. 15:39-47(1996).
EMBL; S83371; AAD14433.1; -.
 094vc6 varanus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Lagomorpha, Leporidae, Oryctolagus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE FROM N.A.
MEDLINE=96377339; PubMed=8783186;
Metsaranta M., Kujala U.M., Pelliniemi L., Osterman H., Aho H.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 19; DB 6; Length 8; Pred. No. 8.3e+05;
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01-JUL-1997 (TrEMBLrel. 04, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
Pro alpha 1 type III collagen protein (Fragment).
Oryctolagus cuniculus (Rabbit).
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8 AA; 1028 MW; B859C7272EA77371 CRC64;
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Q8W8X4;
01-MAR-2002 (TrEMBLrel: 20, Created)
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098TU5
098TU5
009228
0052062
015901
002032
095TT7
095ZY1
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Q94NB2
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Q9PRJ4
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ID Q8W8;
AC Q8W8;
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Q94vcl varanus rud
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                                                                                                     November 17, 2003, 18:27:09; Search time 27.5 Seconds (without alignments) 84.454 Million cell updates/sec
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(c) 1993 - 2003 Compugen Ltd.
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Gapop 10.0 , Gapext 0.5
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Perfect score:
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Genome Res. 7:142-156(1997).
EMBL; U68389; AAB19199.1; -.
TRANSFAC; T03410; -.
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66.7%;
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nes 3; Conservative
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Matches 2; Conservative
                                                             PRELIMINARY;
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01-MAR-2002 (TrEMBLrel.
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                                                                                                                                                                                                          Human herpesvirus 1.
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                                                                                                                                                                                                                                                                                                              SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                      NCBI TaxID=10298;
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ID Q69473
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Lepidosauria; Squamata; Scleroglossa; Anguimorpha; Varanidae; Varanus.
NCBI_TaxID=169851;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Ast J.C.; "Mitochondrial DNA evidence and evolution in Varanoidea (Squamata)."; Cladistics 17:0-0(2001).
EMBL; AF407521; AAL10116.1; -.
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STRAIN=CC70, and CC117;
MEDLINE=21561594; PubMed=11703875;
Lessios H.A., Garrido M.J., Kessing B.D.;
"Demographic history of Diadema antillarum, a keystone herbivore on Caribbean reefs.";
                                                                                                                         Eukaryota, Metazoa, Echinodermata, Eleutherozoa, Echinozoa,
Echinoidea, Euechinoidea, Diadematacea, Diadematoida, Diadematidae,
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MEDLINE=21323357; PubMed=11430656;
Lessios H.A., Kessing B.D., Pearse J.S.;
Population structure and speciation in tropical seas: global phylogeography of the sea urchin Diadema.";
Evolution 55:955-975(2001).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 18; DB 8; Length 8; Pred. No. 8.3e+05; 1; Mismatches 0; Indels
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EMBL, AY012920; AAL33843.1; -
EMBL; AY012921; AAL33844.1; -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                9 AA; 1174 MW; 2B73173B46DDC2D3 CRC64;
01-MAR-2002 (TrEMBLrel. 20, Last sequence update) 01-MAR-2002 (TrEMBLrel. 20, Last annotation update)
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                                                                                     Diadema mexicanum.
                                                                                                                                                                                                                                  SEQUENCE FROM N.A.
                                                                                                                                                                                   NCBI_TaxID=105359
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Matches
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                                                                                                                                                                                                                                                                                                                                                              Chang Y., Jeang K., Lietman T., Hayward G.S.;
"Structural Organization of the Spliced Immediate-Early Gene Complex that Encodes the Major Acidic Nuclear (IE1) and Transactivator (IE2) Proteins of African Green Monkey Cytomegalovirus.";
J. Biomed. Sci. 2:105-13(1995).
EMBL. U18080; AAA75442.1;
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Amphibia; Batrachia; Anura; Mesobatrachia; Pipoidea; Pipidae;
Xenopodinae; Xenopus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDLINE=97202105; PubMed=9049612; Steelman S., Moskow J.J., Muzynski K., North C., Druck T., Montgomery J.C., Huebner K., Daar I.O., Buchberg A.M.; "Identification of a conserved family of Meis1-related homeobox
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                                                                                                                                                                                                 Viruses, dsDNA viruses, no RNA stage, Herpesviridae, Alphaherpesvirinae, Simplexvirus.
                                                                      01-NOV-1996 (TrEMBLrel. 01, Last sequence update) 01-MAR-2002 (TrEMBLrel. 20, Last annotation update) Immediate-early transactivator 110 (Fragment).
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Last annotation update)
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. 8.3e+05;
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9 AA.
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                                                 Created)
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Xenopus laevis (African clawed frog)
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PRT;
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Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Buteleostomi,
Lepidosauria, Squamata, Scleroglossa, Anguimorpha, Varanidae, Varanus.
VCBI_TaxID=169832;
                                                                                                                                                                                                                                                                      Ast J.C.;
"Mitochondrial DNA evidence and evolution in Varanoidea (Squamata).";
[Cladistics 17:0-0(2001).
EMBL; AF407524; AAL10125.1; -.
Mitochondrion.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STRAIN=1, 2, 7, 12, 16, 17, and 110;
Randi E., Pierpaoli M., Beaumont M., Ragni B., Sforzi A.;
Randi E., Pierpaoli M., Beaumont M., Ragni B., Sforzi A.;
Randi E., Pierpaoli M., Beaumont M., Ragni B., Sforzi A.;
and their hybrids using Bayesian clustering methods.";
Submitted (WAR-2001) to the EMBL/GenBank/DDBJ databases.
EMBL; AJ409129; CAC41037.1; --
EMBL; AJ409139; CAC41033.1; --
EMBL; AJ409132; CAC41039.1; --
EMBL; AJ409132; CAC41039.1; --
EMBL; AJ409133; CAC41049.1; --
EMBL; AJ409134; CAC41045.1; --
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Bukaryota, Metazoa, Chordata, Craniata, Vertebrata, Buteleostomi,
Mammalia, Butheria, Carnivora, Fissipedia, Felidae, Felis.
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01-DEC-2001 (TYEMBLrel. 19, Last sequence update)
01-DEC-2001 (TYEMBLrel. 19, Last annotation update)
Cytochrome c oxidase subunit II (Fragment).
                                                                                  Last sequence update)
Last annotation update)
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                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE 8 AA; 992 MW; EFC775A5A36411A6 CRC64;
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                                                                 01-DEC-2001 (TrEMBLrel. 19, Created)
01-DEC-2001 (TrEMBLrel. 19, Last sequence of 01-DEC-2001 (TrEMBLrel. 19, Last annotation Cytochrome c oxidase subunit I (Fragment).
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nes 2; Conservative
                                 PRELIMINARY;
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SEQUENCE FROM N.A.
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                                                 094VB2;
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01-DEC-2001 (
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 RESULT 8
094VB2
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TISSUE=Placenta;
Lee C.-C., Yazdani A., Wehnert M., Bailey J., Couch L., Xiong M.,
Coolbaugh M.I., Chinault C.A., Baldini A., Lindsay B.A., Zhao Z.-Y.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mitochondrion.
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Carnivora, Fissipedia, Felidae, Felis.
                                                                                                                Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
                                                                                                                                                                                                                                                 Caskey C.T.H.;
"Isolation of chromosome-specific genes by reciprocal probing of arrayed cDNAs and cosmid libraries.";
Hum. Mol. Genet. 0:0-0(1995).
EMBL: L32069; AAA73878.1; -
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01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
Cytochrome c oxidase subunit II (Fragment).
                            01-NOV-1996 (TrEMBLrel. 01, Created)
01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                          8 AA; 1068 MW; 0315A37EAB5B0763 CRC64;
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8 AA; 951 MW; 262685BDC5A3733B CRC64;
 8 AA.
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 PRT;
                                                                             (Clone XP15H8A) (Fragment).
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Best Local Similarity 100.
Matches 2; Conservative
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PRELIMINARY;
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                                                                                                  Homo sapiens (Human)
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Best Local Similarity
Matches 2; Conserv
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SEQUENCE 8 AA
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RESULT 10 **094PX6**

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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Lepidosauria; Squamata; Scleroglossa; Anguimorpha; Varanidae; Varanus.
NCBI_TaxID=169830;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STRAIN-Sprague-Dawley; TISSUE-Liver;
MEDLINE=20198203; PubMed=10731662;
Kim K.-Y., Choi I., Kim S.-S.,
"Purification and characterization of a novel inhibitor of the
proliferation of hepatic stellate cells.";
J. Biochem. 127:23-27(2000)
-!-FUNCTION: MAY ACT AS A NEGATIVE EFFECTOR IN THE REGULATION OF THE
HEPATIC STELLATE CELLS (HSC). ALSO INHIBITS THE GROWTH OF BOVINE
ENDOTHELIAL CELLS AND 376 FIBROBLASTS.
-!- SIMILARITY: IDENTICAL TO THE 63-70 AA REGION OF THE RAT ZAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  "Mitochondrial DNA evidence and evolution in Varanoidea (Squamata).";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Rattus. NCBI_TaxID=10116;
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01-0CT-2000 (TrEMBLrel. 15, Last sequence update)
01-MR-2010 (TrEMBLrel. 16, Last annotation update)
38kDa non-arginase growth inhibitory factor (NAGIF) (Fragment).
Rattus norvegicus (Rat).
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Last annotation update)
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                                                  8 AA.
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                                                                                                           Created)
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EMBL, AF407523, AAL10122.1; -.
Mitochondrion.
                                                                                                  01-DEC-2001 (TrEMBLrel. 19, 01-DEC-2001 (TrEMBLrel. 19, 01-DEC-2001 (TrEMBLrel. 19,
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Matches 2; Conservative
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                                                  PRELIMINARY;
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Best Local Similarity
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                                                                                                                                                                                                                                                                                   Mitochondrion
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P82598
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RESULT 12
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Mitochondriion.
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Lepidosauria; Squamata; Scleroglossa; Anguimorpha; Varanidae; Varanus.
NCBI_TaxID=169831;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ast J.C.; "Microchondrial DNA evidence and evolution in Varanoidea (Squamata)."; Cladistics 17:0-0(2001). EMBL; AF407526; AAL10130.1; -. Mitochondrion.
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Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Carnivora, Fissipedia, Felidae, Felis.
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                                                                                                                                                              01-DEC-2001 (TrEMBLrel. 19, Created)
01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
Cytochrome c oxidase subunit II (Fragment).
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8 AA; 951 MW; 262685BDC5A3733B CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                            NCBI_TaxID=61377;
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                                                                                                                                                                                                                  "Nucleotide sequences at the 5'-termini of the alfalfa mosaic virus RNAs and the intercistronic junction in RNA 3.";
Nucleot Acids Res. 8:5635-5647(1980).
EMBL: V00047; CAA23416.1; -
NON_TER
                                                                         Alfalfa mosaic virus.
Viruses; ssRNA positive-strand viruses, no DNA stage; Bromoviridae;
                                                                                                                                                       SEQUENCE FROM N.A.
MEDILINE=81124289; PubMed=6927843;
Koper-Zwarthoff B.C., Brederode F.T.M., Veeneman G., van Boom J.H.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Eukaryota, Metazoa; Chordata, Craniata, Vertebrata, Euteleostomi, Mammalia; Eutheria; Primates; Catarrhini, Hominidae; Homo. NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE FROM N.A.
MEDLINE=21539745; PubMed=11682035;
Kagaya M., Kaneko S., Ohno H., Innarra K., Kobayashi K.;
Kagaya M., Kaneko S., Ohno H., Innarra K., Kobayashi K.;
"Cloning and characterization of the S. flanking region of human cytokeratin 19 gene in human cholangiocarcinoma cell line.";
C. Hepptol. 35:504-511(2001).
EMBL; AB045973; BAB40770.1; -.
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             01-NOV-1996 (TrEMBLrel. 01, Created)
01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
Putative ORF (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                01-JUN-2001 (TrEMBLrel. 17, Created)
01-JUN-2001 (TrEMBLrel. 17, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
Cytokeratin 19 (Fragment).
                                                                                                                                                                                                                                                                                                       SEQUENCE 8 AA; 917 MW; 69D40B0775A365B8 CRC64;
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                                                                                                                         NCBI_TaxID=12321;
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SEQUENCE
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Q9BYF9
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Search completed: November 17, 2003, 18:31:19 Job time : 29.5 secs

3 SYSYR 7

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Gonadotrophin rele Gonadotrophin rele Gonadotrophin rele Core polypeptide f DP178-like/DP107-l

Seabream gonadotro Luteinising hormon Luteinising hormon

Immunogenic LHRH(3 LHRH, peptide fragm Human LHRH peptide Sequence of lutein Gonadotropin relea

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Reproduction stimu Des-G1y10, (D-Ala6 Peptide for stimul Luteinising hormon Luteinising hormon Syntheric peptide Nonapeptide ethyla Water insoluble li

Peptide #3 generat Luteinising Hormon Luteinising Hormon Gonadoliberin anal

LH-RH analogue 3

Luteinising hormon

Title: Perfect score:

Sequence:

OM protein

Run on:

Scoring table:

Searched:

Minimum DB Maximum DB

Database

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LHRH; immune response; luteinising hormone releasing hormone; DT; diphtheria toxoid; castrating; oestrus cycling; aggression; breast; sexual activity; organoleptic; livestock; cell growth; malignant; prostate; ovarian; oncofoetal; hyperplastic; pregnancy;
                                                                                                                                                                                                                                                                                                                                                                                                                                              ALIGNMENTS
                                                                                                             AAW94892
AAB15364
AAP60174
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AAB90973
AAB37133
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AAU76984
ABP96547
AAW47843
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AAW03070
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AAR89932
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AAP20240
AAP50837
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AAP10415
AAP10007
AAP20405
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAW94891 standard; peptide; 9 AA.
98WO-AU00532
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(CSTC-) CST TLD
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AAW94891;
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AAW94891
LHRH peptide fragm
Human LHRH peptide
Amino acid sequenc
Luteinising hormon
GnRH peptide. Pet
Luteinising hormon
Gonadotrophin rele
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                                                                          November 17, 2003, 18:23:44; Search time 34 Seconds (without alignments) 42.016 Million cell updates/sec
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Compugen Ltd
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         GenCore version (c) 1993 - 2003
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Maximum Match 100%
Listing first 45 summaries
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AAB08104
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Gapop 10.0 , Gapext 0.5
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58
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                                                                                                                                                                                                                                               response to luteinising hormone releasing hormone (LHRH). The composition comprises a LHRH-diphtheria toxoid (DT) conjugate adsorbed to an ionic polysaccharide. The LHRH-DT compositions can be used for eliciting an immune response to LHRH, for castrating an animal, for regulating osetrus cycling in a female animal or for inhibiting characteristics induced by the sexual maturation of an animal, e.g. aggression or sexual activity. They can also be used for achieving production gains in livestock, e.g. reduction or elimination of unwanted cygnoclopic characteristics from the meat of livestock. They can also be used for inhibiting the growth of cells which are regulated directly or indirectly by LHRH, e.g. malignant breast cells, malignant prostate cells, malignant ovarian cells, malignant oracines and seed for inhibiting they can also be used for down-regulating the libido of an animal. They can also be used for inhibiting pregnancy, prostate cells. They can also be used for inhibiting pregnancy, prostate cells animal. They can also be used for inhibiting pregnancy, prostate cells animal. They can also be used for inhibiting pregnancy, prostate cells animal. They can also be used for inhibiting pregnancy, prostate compositions induce a more effective immune response against LHRH than the carrier of the LHRH carrier.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                against LHRH results in prevention of the release of the hormones LH and FSH from the anterior pituitary. Sequences AAW94890-93 are peptide derivatives of LHRH.
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                                                                                                                                                                                                                 The invention relates immunogenic composition for eliciting an immune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
New immunogenic leutenising hormone releasing hormone compositions comprise LHRH conjugated to diphtheria toxoid and adsorbed to an ionic polysaccharide, used to inhibit reproductive function in
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gonadotrophin releasing hormone; fertility control; cancer;
endometriosis; prostate enlargement.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    100.0%; Score 58; DB 20; Length 9; 100.0%; Pred. No. 9.3e+05;
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                                                                                                                                                               3; Page 30; 41pp; English.
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Matches 9; Conservative
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                                                                The present sequence is a peptide fragment of human luteinising hormone releasing hormone (also known as LHRH, GnRH and gonadotrophin releasing hormone). It was used to demonstrate the novel adjuvant of the invention, which has lower reactogenicity then previous compositions. Vaccination of humans and animals against LHRH can be used as a method of fertility control, as well as enabling the control and treatment of fertility the reproductive organs, such as testicular, breast, prostate and ovarian cancers, prostate enlargement and endomerriosis. The composition of the invention contains an anionic macromolecule and a saponin component, the latter of which is an immunostimulant, and it can also be used with other immunogens including soluble protein antigens, peptide haptens conjugated to a carrier protein and whole viruses.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present sequence represents luteinising hormone releasing hormone (LHRH). It is used in vaccines with T helper cell epitopes AAB08076-808101, derived from canine distemper virus (CDV). Compositions comprising these T cell helper epitopes are useful for inducing an immune response in an animal. The epitopes are useful as components of animal, in particular, canine vaccines, either simply as synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel T helper cell epitopes derived from canine distemper virus useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Amino acid sequence of truncated luteinising hormone releasing hormone.
                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 T helper cell epitope; CDV; immune response; canine vaccine; luteinising hormone releasing hormone; LHRH.
                                                                                                                                                                                                                                                                                                                                                                                                                        0; Indels
                                                                                                                                                                                                                                                                                                                                                                                Length
                                                                                                                                                                                                                                                                                                                                                                              100.0%; Score 58; DB 21; 100.0%; Pred. No. 9.3e+05;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        for preparation of canine vaccines
                                     Disclosure; Page 50; 53pp; English.
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99AU-0002013.
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immunostimulating complex
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nes 9; Conserv
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                                                                                                                                                                                                                                                                                                                                           9 AA;
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                                                                                                                                                                                                                                                                                                                                             Sequence
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                                                                                                                                                                                                                                                                                                                                                                                Query Match
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Length 9;

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Query Match
Best Local Similarity
Matches 9; Conserv
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                                                                                                                                                                                                                                                                                                       Key
Modified-site
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03-JUN-1999;
                                                                                                                                                                                                         GnRH peptide
                                                                                                                                                                                 26-MAR-2001
                                                                                                                                                                                                                                                                                                                                                                              14-DEC-2000.
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                                                                                                                                                          AAB59836;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention describes a modified therapeutic peptide (I) comprising a therapeutically active amino acid region (III) and a reactive group (II) (e.g. succinimidy) and maleimido groups) attached to a less therapeutically active amino acid region (IV), which covalently bonds with amino/hydroxyl/thiol groups on blood components to form a peptidase stabilised therapeutic peptide composed of 3-50 amino acids. (I) are useful for modifying therapeutic peptides e.g. hormones, growth factors and neurotransmitters, to protect them from peptidase activity in vivo for the treatment of various disorders. Endogenous therapeutic peptides are not suitable as drug candidates as they require frequent administration due to rapid degradation by peptidases in the body. Modifying and attaching therapeutic peptides to albumin prevents or reduces the action of peptidases to increase length of activity (half life) and specificity as bonding to large molecules decreases the action of peptidases to increase length of activity (half intracellular uptase and interference with physiological processes. ABB90829 to AAB92441 represent invention.
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                                                                                                                                                                                                                                                                    Luteinising hormone releasing hormone (LH-RH) related peptide SEQ ID:153.
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                                                                                    Gaps
                                                                                                                                                                                                                                                                                         Protection; endogenous therapeutic peptide; peptidase; conjugation; blood component; modification; succinimidyl; maleimido group; amino; hydroxyl; thiol; hormone; growth factor; neurotransmitter.
peptide based vaccines and as additions to vaccines containing more
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                                                            Length 9;
                                                                                 Indels
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                                                           Score 58; DB 21;
Pred. No. 9.3e+05;
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                                                                                   0; Mismatches
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                                                                                                                                                                                           AAB90979 standard; Peptide; 9 AA.
                                                         100.0%;
100.0%;
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99US-0153406.
99US-0159783.
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                                                                                                                                                                                                                                           22-JUN-2001 (first entry)
                                                        Query Match
Best Local Similarity 100.
Matches 9; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Bridon DP, Ezrin AM,
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                                                                                                                        1 HWSYGLRPG 9
                                                                                                         1 HWSYGLRPG 9
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               complex antigens
                                    9 AA;
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10-SEP-1999;
15-OCT-1999;
                                                                                                                                                                                                                                                                                                                                            Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                        Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GnRH-III; autoimmune disease; transplant rejection; retroviral disease; graft-versus-host-disease; lymphoproliferative disease; gonadotropin-releasing hormone.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Novel product comprising proliferatively active moiety linked to genetic material, useful as vectors for protected nucleic acid material and as mitogen to stimulate proliferation of target cell -
                                                           Gaps
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                                                           Indels
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100.0%; Pred. No. 9.3e+05;
ive 0; Mismatches 0;
100.0%; Score 58; DB 22;
100.0%; Pred. No. 9.3e+05;
tive 0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                         AAB59836 standard; Peptide; 9 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
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Matches 9; Conservative
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HWSYGLRPG
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AAB90983

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The present invention describes a pharmaceurical composition (1) to the present invention describes a pharmaceurical composition (1) to treating breast cancer, endometriosis, myometrium tumour, Alzhaimer's disease, circulatory system disorders, menopausal disorders, irregular confectorations, cancer metastasis, premenstrual syndrome, muscular distress or osteopathies due to calcium/phosphorus imbalance. (1) comprises a consider of considering the selective cestrogen control of considering the selective cestrogen control of considering muscular and osteopathic neuroprotective, cardiovascular, andocrine, muscular and osteopathic activities. (1) can be used as GnRH agonists used in combination with controlive oestrogen receptor modulators, selective androgen receptor modulators, sex hormone synthesis inhibitors, lyase inhibitors, controlive oestrogen receptor controlive oestrogen receptor modulators, bone metabolism regulators, receptor receptor-type tyrosine kinase inhibitors, bone metabolism regulators, antagonists for treating and preventing endometriosis, myometrium tumour, cirregular periods, cancer metastasis, premenstrual syndrome, muscular distress, osteopathies due to calcium/phosphorus imbalance, prostatomegally or prostate cancer or breast cancer or their reccurrence conferences, cancer activity and/or reducing side effects. The present sequence cor represents a GnRH agonist related peptide which is given in the present
                                                                                                  Gonadotrophin releasing hormone agonist; GnRH agonist; combination drug; pharmaceutical; breast cancer; endometriosis; myometrium tumour; Alzheimer's disease; circulatory system disorder; menopausal disorder; irregular period; cancer metastasis; premenstrual syndrome; osteopathy; muscular distress; calcum/phosphorus imbalance; SERM; gynaecological; selective oestrogen receptor modulator; cytostatic; nootropic; muscular; neuroprotective; cardiovascular; endocrine; osteopathic; prostatomegally;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Pharmaceutical composition e.g. for breast cancer comprises gonadotrophin releasing hormone agonist and selective estrogen receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  present invention describes a pharmaceutical composition (1) for
                                                 Gonadotrophin releasing hormone agonist gonadrelin peptide.
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Pred. No. 9.3e+05;
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100.0%; Pre-
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27-MAY-2003 (first entry)
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                                                                                                                                                                                                                                                                                               prostate cancer.
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                                                                                                                                                                                                                                                                                                                                                 Synthetic.
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                                                                                                                                                                                       Luteinising hormone releasing hormone (LH-RH) related peptide SEQ ID:157.
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                                                                                                                                                                                                                                      Protection; endogenous therapeutic peptide; peptidase; conjugation; blood component; modification; succinimidyl; maleimido group; amino; hydroxyl; thiol; hormone; growth factor; neurotransmitter.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              intracellular uptake, and interference with physiological processes. AAB90829 to AAB92441 represent peptides which can be used in the exemplification of the present invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 55, DB 22; Length 9; Pred. No. 9.3e+05; 1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Holmes DL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure; Page 241; 733pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Milner PG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABP96550 standard; peptide; 8 AA.
                           AAB90983 standard; Peptide; 9 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    99US-0134406.
99US-0153406.
99US-0159783.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               94.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                17-MAY-2000; 2000WO-US13576
                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    8; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (CONJ-) CONJUCHEM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Ezrin AM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2001-112059/12.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
Matches 8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              9 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                           WO200069900-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    17-MAY-1999;
10-SEP-1999;
15-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                 Homo sapiens
                                                                                                                                   22-JUN-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               23-NOV-2000
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                                                                                                                                                                                                                                                                                                                                                                            Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABP96550;
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                                                                               AAB90983;
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Gaps

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RESULT 7

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ABP96550

Length 8; Indels

Mammalia.

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caused by inadequate secretion of endogenous parathormone (PTH) which contains, apart from an acceptable carrier, gonadoliberin or agonists at least as strongly active as gonadoliberin. For parenteral use these provide 0.5-5 micrograms gonadoliberin per unit dose, and for application to mucosa (intranasally) 10-200 micrograms (Updated on 03-0CT-2002 to add missing OS field.)
                                                                                                                                                                                                                      /label= bonded to -NHCH3, -NH-CH2-CH3, -NH-CH2CH2CH3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Compsn. contg. gonadoliberin or its analogues - for treating parathormone deficiency states, e.g. hypocalcaemic conditions or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The inventors claim a composn. for treating metabolic disorders
                                                                          Gonadoliberin agonist, follitropin release, lutropin release, parathormone, hypertension, therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gonadotropin releasing hormone; GnRH; sbGnRH; gonadoliberin; spawning; ovulation; fish farming; transgenic fish; seabream
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 9;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 6; Le
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Seabream gonadotropin releasing hormone sbGnRH-I.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    89.7%; Score 52; DB
100.0%; Pred. No. 9.3
:ive 0; Mismatches
                                            Sequence of gonadoliberin analogue IIc.
                                                                                                                                                         Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; Page 6-7; 17pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAR97760 standard; Peptide; 9 AA.
                                                                                                                                                                                       /label= pyroGlu
                                                                                                                                                                                                                                                                                                                     83DE-3332329
                                                                                                                                                                                                                                                                                                                                                    83DE-3332329
 (updated)
(first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     8; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                 Konig W, Neubauer H;
                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1985-081717/14.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Φ
                                                                                                                                                                                                                                                                                                                                                                                  (FARH ) HOECHST AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             HWSYGLRP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         9 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sparus aurata,
                                                                                                                                                                          Modified-site
                                                                                                                                                                                                        Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               hypertension
                                                                                                                                                                                                                                                                                                                   08-SEP-1983;
                                                                                                                                                                                                                                                                                                                                                    08-SEP-1983;
 03-OCT-2002
29-NOV-1991
                                                                                                                           Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO9617619-A1
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                                                                                                                                                                                                                                                                                     28-MAR-1985.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   요
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 composition is used to treat prostate adenocardinoma, benign hypertrophy of the prostate, endometriosis, dysmenorrhea, hirsutism, hornome-dependent mammary tumours, for treatment or prevention of precocious puberty, delaying the onset of puberty and for treating see AAP10411-P10418.

(Updated on 10-VMR-2002 to add missing PI field.)

(Updated on 10-MRR-2003 to correct PA field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                       LHRH; Follicle Stimulating Factor; FSH; acne; hirsutism; dysmenorrhea; precocious puberty; endometriosis; prostate cancer; benign prostate hypertrophy; mammary tumour.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LH-RH, liberating factor for LH and FSH, and its agonists compsn. - used to treat prostate adenocarcinoma, benign hypertrophy of the prostate, hirsutism, acne, etc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ;
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       composition is claimed containing LHRH or its analogues. The
                                                                                                                                                                                                                                                                                                                                "Pro-NH-(CH2)n-CH3 (n=0-2),
Pro-NH-(CH2)2-OH or protected by
pyrrolidino or morpholino gp."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 9;
                                                                                                           Luteinising Hormone Releasing Hormone analogue #3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               89.7%; Score 52; DB 2; Le. 100.0%; Pred. No. 9.3e+05; ive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                    /note= "pyroglutamic acid"
                                                                                                                                                                                                                                                      Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 1(d); Page 15; 27pp; French.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAP50568 standard; Protein; 9 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1981-23409D/14 (23409D).
                                                                                                                                                                                                                                                                                    label= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               79FR-0023545
                                                                                                                                                                                                                                                                                                                                                                                                                                                               83BE-0468932
                                                           (updated)
(first entry)
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                                                                                                                                                                                                                                                                                                                                     /note=
                             (updated)
(updated)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (ROUS ) ROUSSEL-UCLAF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Labrie F, Raynaud J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 HWSYGLRP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  9 AA;
                                                                                                                                                                                                                                                      Key
Modified-site
                                                                                                                                                                                                                                                                                                                   Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                               23-FEB-1983;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             21-SEP-1979;
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                             25-MAR-2003
                                              10-MAR-2003
                                                           01-JUL-2002
17-DEC-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                19-MAR-1981
                                                                                                                                                                                                                                                                                                                                                                                                 BE885308-A
                                                                                                                                                                                                                        Synthetic.
AAP10414;
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Gaps

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Sequence

Query Match Best Local

Matches

AAP50568

AAPSOS68 ID AAPS XX AC AAPS

RESULT 9

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Disclosure; Page 238; 733pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2001-112059/12.
                                                                                                                                                                                                                                                       1 HWSYGLRP 8
                                                                                                                                                                                                                   Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                         HWSYGLRP
                                                                                                                                                                                               9 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                      sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       17-MAY-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                10-SEP-1999;
15-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   23-NOV-2000.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DP,
                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic
                                                                                                                                                                                                                                                                                                                                          AAB90986;
                                                                                                                                                                                                Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Bridon
                                                                                                                                                                                                                                      Matches
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                                                                                                                                                                                                                                                                                    ö
                                                                                                                                                                                                                                                                                                                                                                                                                            Luteinising hormone releasing hormone (LH-RH) related peptide SEQ ID:146.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Modifying and attaching therapeutic peptides to albumin prevents peptidase degradation, useful for increasing length of in vivo activity
                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                            Protection; endogenous therapeutic peptide; peptidase; conjugation; blood component; modification; succinimidyl; maleimido group; amino; hydroxyl; thiol; hormone; growth factor; neurotransmitter.
                                                                                                                                                                    A seabream pitutary extract was subjected to HPLC. A fraction showing gonadotropin releasing hormone (GRRH) activity was subjected to pyroglutamyl aminopeptidase digestion and the peptide obtd. (AAR97760) was sequenced. Based on the novelty of the Ser residue at position 8, this peptide was named sbGnRH-I (see also AAR97757). sbGnRH is useful for controlling reproduction in fish.
                                                                                                                      Novel seabream gonadotropin-releasing hormone and its analogues useful for controlling gonadal development and spawning in fish
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Thibaudeau K;
                                                                                  Zohar Y;
                                                                                                                                                                                                                                                               Score 52; DB 17; Length 9;
Pred. No. 9.3e+05;
0; Mismatches 1; Indels
                                                                                  Rivier JEF, Sherwood NM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Holmes DL,
                                                      (UYMA-) UNIV MARYLAND BIOTECHNOLOGY INST.
(UYVI-) UNIV VICTORIA INNOVATION & DEV CORP.
                                                                                                                                                  Disclosure; Page 24; 63pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Milner PG,
                                                                                                                                                                                                                                                                                                                                                                    AAB90972 standard; Peptide; 9 AA.
                                                                                                                                                                                                                                                               89.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   17-MAY-2000; 2000WO-US13576.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     99US-0134406.
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                  95WO-US15886
                                                                                                                                                                                                                                                                                                                                                                                                         22-JUN-2001 (first entry)
                                                                                                                                                                                                                                                                                   8; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (CONJ-) CONJUCHEM INC.
                                                                                  Gothilf Y, Powell J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Bridon DP, Ezrin AM,
                                                                                                                                                                                                                                                                                                    1 HWSYGLRPG 9
                                                                                                    WPI; 1996-286922/29.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2001-112059/12.
                                                                                                                                                                                                                                                                                                                HWSYGLSPG
                                                                                                                                                                                                                                                                          Local Similarity
                                                                                                                                                                                                                                              9 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO200069900-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      17-MAY-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens
Synthetic.
                  04-DEC-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             10-SEP-1999;
15-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  23-NOV-2000
13-JUN-1996
                                                                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                                                   RESULT 11
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comprising a thereby description described among a comprising a thereby described by the comprising a thereby described by the comprising a thereby described by the comprising a less therapeutically active amino acid region (III) and a less therapeutically active amino acid region (IV), which covalently bonds with amino/hydroxyl/thiol groups on blood components to form a peptidase stabilised therapeutic peptide components to form a peptidase stabilised therapeutic peptides of 3-50 amino acids. (I) are useful for modifying therapeutic peptides of 3-50 amino acids. (I) are useful for modifying therapeutic peptides of hormones, growth factors and neurotransmitters, to protect them from peptidase activity in vivo for the treatment of various disorders. Endogenous therapeutic peptides as they require frequent administration due to rapid degradation by peptidases in the body.

Modifying and attaching therapeutic peptides to albumin prevents or reduces the action of peptidases to increase length of activity (half life) and specificity as bonding to large molecules decreases intracellular uptake and interference with physiological processes.

MAB90891 to AAB92441 represent peptides which can be used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        .
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Modifying and attaching therapeutic peptides to albumin prevents peptidase degradation, useful for increasing length of in vivo activity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Protection; endogenous therapeutic peptide; peptidase; conjugation; blood component; modification; succinimidyl; maleimido group; amino; hydroxyl; thiol; hormone; growth factor; neurotransmitter.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention describes a modified therapeutic peptide (I) comprising a therapeutically active amino acid region (III) and a
present invention describes a modified therapeutic peptide (I)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ..
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    89.7%; Score 52; DB 22; Length 9; 100.0%; Pred. No. 9.3e+05; rive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Holmes DL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; Page 242; 733pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Milner PG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             89.7.,
100.0%; PL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAB90986 standard; Peptide; 9 AA.
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99US-0153406.
99US-0159783.
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reactive group (II) (e.g. succinimidyl and maleimido groups) attached to a less therapeutically active amino acid region (IV), which covalently bonds with amino/haroxyl/thiol groups on blood components to form a peptidase stabilised therapeutic peptide composed of 3-50 amino acids.

(I) are useful for modifying therapeutic peptides e.g. hormones, growth factors and neurotransmitters, to protect them from peptidase activity; in vivo for the treatment of various disorders. Endogenous therapeutic peptides are not suitable as drug candidates as they require frequent oppidas are not suitable as drug candidates in the body.

Modifying and attaching therapeutic peptides to albumin prevents or reduces the action of peptidases to increase length of activity (half life) and specificity as bonding to large molecules decreases in the action of peptidases to increase length of activity (half intracellular uptake and interference with physiological processes.

ARB90829 to AAB92441 represent poptides which can be used in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                    ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gonadotrophin releasing hormone agonist; GnRH agonist; combination drug; pharmaceutical; breast cancer; endometriosis; myometrium tumour; Alzheimer's disease; circulatory system disorder; menopausal disorder; irregular period; cancer metastasis; premenstrual syndrome; osteopathy; muscular distress; calcium/phosphorus imbalance; SERM; gynaecological; selective oestrogen receptor modulator; cytostatic; nootropic; muscular; neuroprotective; cardiovascular; endocrine; osteopathic; prostatomegally;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present invention describes a pharmaceutical composition (1) for treating breast cancer, endometriosis, myometrium tumour, Alaheimer's disease, circulatory system discorders, menopausal disorders, irregular periods, cancer metastasis, premenstrual syndrome, muscular distress or osteopathies due to calcium/phosphorus imbalance. (1) comprises a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Pharmaceutical composition e.g. for breast cancer comprises gonadotrophin releasing hormone agonist and selective estrogen receptor
                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                    .
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gonadotrophin releasing hormone agonist nafarelin peptide.
                                                                                                                                                                                                                                                                                                                                                                            Length 9;
                                                                                                                                                                                                                                                                                                                                                                         Score 52; DB 22; Length 9;
Pred. No. 9.3e+05;
0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure; Page 8; 73pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABP96548 standard; peptide; 9 AA
                                                                                                                                                                                                                                                                                                                                                                            88.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         10-AUG-2001; 2001JP-0244616.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   08-AUG-2002; 2002WO-JP08130.
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                                                                                                                                                                                                                                                                                                                                                      Ouery Match
Best Local Similarity 88.5
امات 8; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 HWSYSLRPG 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 HWSYGLRPG 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2003-300573/29.
                                                                                                                                                                                                                                                                                                                                     9 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO2003015820-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   prostate cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        27-MAY-2003
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                                                                                                                                                                                                                                                                                                                                     Sequence
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receptor modulator (SERM). (I) has cytostatic, gynaecological, nootropic, neuroprotective, cardiovascular, endocrine, muscular and osteopathic neuroprotective, cardiovascular, endocrine, muscular and osteopathic cartivities. (I) can be used as GRRH agonists used in combination with selective cestrogen receptor modulators, selective androgen receptor modulators, selective androgen receptor captures is inhibitors, lyase inhibitors, creeptor-type tyrosine kinase inhibitors, bone metabolism regulators, immunotherapy drugs, cytokine/chemokine inhibitors or endothelin receptor antagonists for treating and preventing endometriosis, myometrium tumour, all zhaimer's disease, circulatory system disorders, menopausal disorders, irregular periods, cancer metastasis, premenstrual syndrome, muscular distress, osteopathies due to calcium/phosphorus imbalance, prostatomegally or prostate cancer or breast cancer or their reoccurrence or metastasis. Combination enhances the quality of life by enhancing GRRH agonist activity and/or reducing side effects. The present sequence represents a GRRH agonist related peptide which is given in the present
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gonadotrophin releasing hormone (GnRH) agonist and a selective oestrogen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Agent for preventing post-operative recurrence of premenopausal breast cancer contains GRRH agonists or antagonists -
                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention describes an agent containing gonadotrophin releasing hormone (GnRH) agonists or antagonists. GnRH agonist and antagonists have cytostatic activity. The agent can be used for preventing post-operative recurrence of premenopausal breast cancer. The agents have no serious side effects with sustained drug effect over a long period without frequent administration. The present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /note= "Pro is C-terminally modified with -NHCH2CH3"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              releasing hormone agonist; GnRH agonist; cancer; breast cancer; cytostatic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gonadotrophin releasing hormone agonist peptide gonadrelin.
                                                                                                                                                                                                                                                                                                                                                                                       Score 52; DB 24; Length 9; Pred. No. 9.3e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                           1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                             0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure; Page 7; 39pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABP96021 standard; peptide; 9 AA.
                                                                                                                                                                                                                                                                                                                                                                                     89.78;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              25-APR-2001; 2001JP-0128032.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (TAKE ) TAKEDA CHEM IND LTD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     02-MAY-2003 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                               8; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2003-148286/14.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Igari Y, Kusaka M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 HWSYALRPG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 HWSYGLRPG
                                                                                                                                                                                                                                                                                                                                                                                                           Sest Local Similarity
                                                                                                                                                                                                                                                                                                                                                    9 AA;
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premenopausal
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                                                                                                                                                                                                                                                                                                              invention
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                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
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                                                                                                                                                                                                                                                                                                                                                                 Gonadotrophin releasing hormone analogue; neurotoxin; prostate cancer; endocrine disorder; gonadotrophin related illness; endometrial cancer; pancreatic cancer; breast cancer; endometriosis; precocious puberty; GRRH-A; therapy.
                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New agent comprising a light chain and a (modified) heavy chain of botulinum, butyricum, or tetani toxin, useful for treating a gonadotrophin related illness, e.g. breast, prostate pancreatic or endometrial cancer, or endometriosis
sequence represents an example of a GnRH agonist, from the present
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 9;
                                                                            Length 9;
                                                                                                         0; Indels
                                                                                                                                                                                                                                                                                                                                         Gonadotrophin releasing hormone analogue (GnRH-A) #3.
                                                                           DB 24; L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        y Match 89.7%; Score 52; DB 24; I Local Similarity 100.0%; Pred. No. 9.3e+05; Hes 8; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /note= "C-terminal ethylamide"
                                                                         Query Match 89.7%; Score 52; DB Best Local Similarity 100.0%; Pred. No. 9.3 Matches 8; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                'note= "Pyroglutamic acid"
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                                                                                                                                                                                                                                                 AAE29840 standard; peptide; 9 AA.
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                                                                                                                                                                                                                                                                                                             (first entry)
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2 HWSYGLRP 9
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Modified-site
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                 invention.
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Sequence 204, App
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Sequence 442, App
Sequence 16, App
Sequence 16, Appl
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Sequence 13, Appl
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GenCore version 5.1.6
(c) 1993 - 2003 Compugen Ltd.
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US-09-192-81-117

US-09-191-393-72

US-09-791-393-72

US-09-791-393-72

US-09-791-389-72

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US-10-062-257-16

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16 25 43.1 7 15 US-10-254-466A-239 Sequence 239, App 1 18 25 43.1 9 14 US-10-109-31-6 Sequence 10, Appli 20 25 43.1 9 14 US-10-109-31-6 Sequence 8, Appli 20 25 43.1 9 14 US-10-109-31-10 Sequence 10, Appli 21 25 43.1 9 14 US-10-109-31-12 Sequence 10, Appli 22 43.1 9 14 US-10-109-31-12 Sequence 11, Appli 22 43.1 9 14 US-10-109-31-14 Sequence 11, Appli 23 25 43.1 9 14 US-10-109-31-16 Sequence 14, Appli 24 43.1 9 14 US-10-109-31-20 Sequence 16, Appli 26 25 43.1 9 14 US-10-109-31-20 Sequence 20, Appli 26 25 43.1 9 14 US-10-109-31-20 Sequence 20, Appli 27 43.1 9 14 US-10-109-31-20 Sequence 21, Appli 37 25 43.1 9 14 US-10-109-31-20 Sequence 21, Appli 37 25 43.1 9 14 US-10-109-31-20 Sequence 21, Appli 37 25 43.1 9 14 US-10-109-31-20 Sequence 21, Appli 37 24 41.4 8 12 US-10-109-31-20 Sequence 17, Appli 38 23 39.7 9 12 US-10-139-709-17 Sequence 17, Appli 39.7 9 12 US-10-139-709-17 Sequence 17, Appli 39.7 9 12 US-10-139-709-17 Sequence 17, Appli 39.7 9 12 US-10-189-35-24 Sequence 16, Appli 39.7 9 12 US-10-189-35-24 Sequence 16, Appli 41 22 37.9 6 10 US-09-81-66 Sequence 15, Appli 42 22 37.9 7 12 US-10-23-55-84 Sequence 15, Appli 44 22 37.9 7 12 US-10-23-555-4 Sequence 15, Appli 44 22 37.9 7 12 US-10-23-555-4 Sequence 15, Appli 44 22 37.9 7 12 US-10-23-555-4 Sequence 15, Appli 44 22 37.9 7 12 US-10-23-555-4 Sequence 15, Appli 44 22 37.9 7 12 US-10-23-555-4 Sequence 15, Appli 44 22 37.9 7 12 US-10-23-555-4 Sequence 15, Appli 45 37.9 7 12 US-10-23-23-33 Sequence 15, Appli 45 37.9 8 9 US-10-23-23-33 Sequence 15, Appli
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ALIGNMENTS

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US-09-746-945-2

Sequence 2, Application US/09746945

Sequence 2, Application US/09746945

Parent No. US20020146779A1

GENERAL INFORMATION:

APPLICANT: MILLAR, Alan R

APPLICANT: MILLAR, Alan R

APPLICANT: MILLAR, Alan R

APPLICANT: MCKEE, Colin M

APPLICANT: PLANCATOR CONTINGHAM, IN MCKEE, Colin M

APPLICANT: PLANCATOR UNMER: US/09/746,945

TITLE OF INVENTION: Methods

FILE REFERENCE: P21083WO

CURRENT FAILING DATE: 1998-06-26

PRIOR PILING DATE: 1998-06-26

PRIOR PILING DATE: 1998-06-28

PRIOR FILING DATE: 1998-08-28

PRIOR FILING DATE: 1998-08-28

PRIOR FILING DATE: 1998-06-26

PRIOR FILING DATE: 1998-06-28

NUMBER OF SEQ ID NOS: 14

SEQ ID NO 2

TEMPERATURE:

CORGANISM: Artificial Sequence

FEATURE:

OUGANISM: Artificial Sequence

FEATURE:

OTHER INFORMATION: Protein N-terminal

US-09-746-945-2

Matches 7; Conservative 0; Mismatches 0; Indels 0; Gaps

ON 1 HWSYGLR 9

HWSYGLR 9

HWSYGLR 9

HWSYGLR 9

HWSYGLR 9
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Sequence 4, Application US/09265606 Patent No. US20020034789A1 GENERAL INFORMATION:
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3Y: linear
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es 4; Conserv
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US-09-192-854-117
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Sequence 1000, Application US/10351641

Publication No. US20030186874A1

GENERAL INFORMATION:
APPLICANT: Barney, S.
APPLICANT: Guthrie, K.
APPLICANT: Meruka, G.
APPLICANT: Meruka, G.
APPLICANT: Meruka, G.
APPLICANT: Lambert, D.
TITLE OF INVENTION: HYBRID POLYBEPTIDES WITH ENHANCED PHARMACOKINETIC
TITLE OF INVENTION: HYBRID POLYBEPTIDES WITH ENHANCED PHARMACOKINETIC
TITLE OF INVENTION: HYBRE: US/10/351,641
CURRENT APPLICATION NUMBER: US/10/350,641
PRIOR PRILING DATE: 1999-07-09
PRIOR PILING DATE: 1999-05-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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Sequence 13, Application US/10016283

Sequence 13, Application US/10016283

Publication No. US2020164702A1

GENERAL INFORMATION:

APPLICANT: Valenzuela et al., David M.

TITLE OF INVENTION: NOVEL TYROSINE KINASE RECEPTORS AND LIGANDS

FILE REFERENCE: REG195-B-PCT-US

CURRENT APPLICATION NUMBER: US/10/016,283

CURRENT FILING DATE: 1990-10

PRIOR PILING DATE: 1990-10

PRIOR FILING DATE: 1996-12

PRIOR FILING DATE: 1996-12

NUMBER OF SEQ ID NOS: 36

SOFTWARE: Patentin Ver. 2.0

SEQ ID NO 13
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88.9%; Pred. No. 5.9e+05;
iive 0; Mismatches 0;
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Matches 4; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity 88.9
Matches 8; Conservative
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LENGTH: 8
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RESULT 4

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Gaps
APPLICANT: Zimmermann, Rainer; Park, John E.;
APPLICANT: Rettig, Wolfgang; Old, Lloyd J.
TITLE OF INVENTION: ISOLATED DIMERIC FIBROBLAST ACTIVATION PROTEIN
TITLE OF INVENTION: ALPHA, AND USES THEREOF
CORRESPONDENCE ADDRESS:
ADDRESSEE: Felfe & Lynch
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 117, Application US/09192854

Patent No. US20020068276A1

GENERAL INFORMATION:

APPLICANT: Winter, Greg

APPLICANT: Winter, Greg

TITLE OF INVENTION: Methods for Selecting Functional Peptides

FILE REPERENCE: 3789/72916

CURRENT APPLICATION NUMBER: US/09/192,854

CURRENT FILING DATE: 1998-11-17

EARLIER APPLICATION NUMBER: 60/066,729

EARLIER FILING DATE: 1997-11-21

NUMBER OF SEQ ID NOS: 2.22

SOUTHARE PATENTIN VET: 2.0
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US-09-265-606-4
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                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.5 inch, 2.0 MB storage COMPUTER: IBM PS/2
OPERATING SYSTEM: PC-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 28; DB 9; Ler
Pred. No. 5.9e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: Hanson, No. US20020034789Alman D. REGISTRATION NUMBER: 30,946
REFERENCE/DOCKET NUMBER: LUD 5330.1
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: Wordperfect
CURRENT APPLICATION DATA: . . APPLICATION NUMBER: US/09/265,606
                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE:
CLASSIFICATION
CLASSIFICATION
PRIOR APPLICATION NUMBER:
APPLICATION NUMBER:
FILING DATE:
APPLICATION NUMBER:
08/230,491
FILING DATE:
20-APRLIC-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        48.3%; Scc.
100.0%; Pre
                                                                                                                                       ADDRESSEE: Felfe & Lynch
STREET: 805 Third Avenue
CITY: New York City
STATE: New York
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (212) 688-9200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEPHONE: (212) 688-920
TELEFAX: (212) 838-3884
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
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us-09-462-089-2.closed.rapb

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RESULT 9

US-09-791-389-72

Sequence 72, Application US/09791389

Sequence 72, Application Wolf9791389

Publication No. US20030032773A1

GENERAL INFORMATION:
APPLICANT: Herath Mudiyanselage Athula Chandrasiri
APPLICANT: Parekh, Rajesh Bhikhu
APPLICANT: Parekh, Rajesh Bhikhu
APPLICANT: Parekh, Rajesh Bhikhu
APPLICANT: Parekh, Ronlef, Christian
APPLICANT: Parekh, Ronlef, Christian
APPLICANT: Tyson, Kerry Louise
TITLE OF INVENTION: Proteins, Genes and Their Use for
TITLE OF INVENTION: and Unipolar Depression
FILE REFERENCE: 2543-1-001 N2
CURRENT APPLICATION NUMBER: US/09/791,389

CURRENT FILING DATE: 2000-02-24

PRIOR APPLICATION NUMBER: GB 0004412.3

PRIOR FILING DATE: 2000-02-24

PRIOR APPLICATION NUMBER: GB 0030050.9

PRIOR FILING DATE: 2000-12-08

PRIOR FILING DATE: 2000-12-08

PRIOR FILING DATE: 2000-12-08

PRIOR FILING DATE: 2000-12-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       j. Sequence 72, Application US/09791393
j. Bublication No. US20030032200A1
j. Sequence 72, Application US/09791393
j. Publication No. US20030032200A1
j. GENERAL INFORMATION:
j. APPLICANT: Herath, Herath Mudiyanselage Athula Chandrasiri
j. APPLICANT: Parekh, Rajesh Bhikhu
j. APPLICANT: Herath, Rajesh Bhikhu
j. APPLICANT: Herath, Rajesh Bhikhu
j. TITLE OF INVENTION: Proteins, Genes and Their Use for
j. TITLE OF INVENTION: Diagnosis and Treatment of Bipolar Affective Disorder
j. TITLE OF INVENTION: Diagnosis and Treatment of Bipolar Affective Disorder
j. TITLE OF INVENTION: DIAGNOSIS and Treatment of Bipolar Affective Disorder
j. TITLE OF INVENTION: DIAGNOSIS and Treatment of Bipolar Affective Disorder
j. CURRENT FILING DATE: 2000-01-02
j. EARLIER PILING DATE: 2000-12-03
j. EARLIER FILING DATE: 2000-12-03
j. EARLIER FILING DATE: 2000-12-03
j. EARLIER FILING DATE: 2000-12-12
j. MUMBER OF SEQ ID NOS: 308
j. SEQ ID NO 72
j. EARLIER FASESO for Windows Version 4.0
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                                                                                                                                                                                                        DB 12; Length 9; 5.9e+05;
                                                                                                                                                                                                                   48.3%; Score 28; DB 100.0%; Pred. No. 5.9 tive 0; Mismatches
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                                                                                                                                                                                                                   Query Match
Best Local Similarity 100.
Matches 5; Conservative
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                                                                          TYPE: PRT
ORGANISM: Homo sapiens
US-09-968-744A-204
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Matches 4; Conserv
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4 GLRPG 8
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US-09-791-393-72
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       SEQ ID NO 204
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                                                LENGTH: 9
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Sequence 204, Application US/09968744A

Publication No. US20030148372A1

GENERAL INFORMATION:

APPLICANT Tomlinson, Ian M

APPLICANT Tomlinson, Ian M

APPLICANT Tomlinson, Ian M

TITLE OF INVENTION: Method to Screen Phage Display Libraries with Different Ligands

TITLE OF INVENTION: Method to Screen Phage Display Libraries with Different Ligands

FILE REFERENCE: 8039/1073

CURRENT APPLICATION NUMBER: US/09/968,744A

PRIOR FILING DATE: 1997-10-20

PRIOR FILING DATE: 1997-11-13

PRIOR FILING DATE: 1997-11-13

PRIOR PELING DATE: 1997-11-21

PRIOR PELING DATE: 1997-11-30

PRIOR FILING DATE: 1998-10-20

PRIOR FILING DATE: 1998-10-20

PRIOR PLUMG DATE: 1998-10-20

PRIOR PLUMG DATE: 1908-1350

PRIOR PLUMG DATE: 1908-1350

WUMBER OF SED ID NOS: 350

SOFTWARE: PatentIn version 3.1
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                                                                                                       48.3%; Score 28; DB 9; Length 9; 100.0%; Pred. No. 5.9e+05; ive 0; Mismatches 0; Indels
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100.0%; Pred. No. 5.9e+05;
iive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 204, Application US/09968561A
Patent No. US20020164642A1
GENERAL INFORMATION:
APPLICANT: Tomlinson, Ian M
APPLICANT: Winter, Gregory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity 100.0
Matches 5; Conservative
                                                                                                    Query Match
Best Local Similarity 100.
Matches 5; Conservative
; ORGANISM: Homo sapiens
US-09-192-854-117
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                , ORGANISM: Homo sapiens
US-09-968-561A-204
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US-09-968-561A-204
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Gaps

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OTHER INFORMATION: Peptide corresponding to rat carbonic anhydrase
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Sequence 8, Application US/10165603

Publication No. US20030021792A1

Publication No. US20030021792A1

Publication No. US20030021792A1

APPLICANT: Stevens, Anthony C.

APPLICANT: Stevens, Anthony C.

TITLE OF INVENTION: PROTEINS

FILE REFERENCE: TPTECH.001A

CURRENT APPLICANT: NUMBER: US/10/165,603

PRIOR PELLING DATE: 2002-06-07

PRIOR APPLICATION NUMBER: 60/297,021

PRIOR PLLING DATE: 2001-06-08

PRIOR PLLING DATE: 2001-06-08

PRIOR PLLING DATE: 2001-06-08

NUMBER OF SEQ ID NOS: 33

SOFTWARE: FREUENCE FREUENCE OF WINDOWS VERSION 4.0
                                                                                                                                                                                                                                                                                                                                                                       Length 9;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Indels
TITLE OF INVENTION: 125P5C8: A TISSUE SPECIFIC PROTEIN TITLE OF INVENTION: HIGHLY EXPRESSED IN VARIOUS CANCERS FLE REPERENCE: 129.39U501 CURRENT APPLICATION NUMBER: US/09/809,638 CURRENT FILING DATE: 2001-03-14 NUMBER OF SEQ ID NOS: 746 SEQ ID NOS: 746 SEQ ID NOS: 746 SEQ ID NO 463 LENGTH: 9
                                                                                                                                                                                                                                                                                                                                                                 Query Match
44.8%; Score 26; DB 11;
Best Local Similarity 60.0%; Pred. No. 5.9e+05;
Matches 3; Conservative 1; Mismatches 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 26; DB 14;
Pred. No. 5.9e+05;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: ITOH, Kyogo
TITLE OF INVENTION: Tumor antigen
FILE REPERENCE: GP00-1017
CURRENT APPLICATION NUMBER: US/10/062,257
CURRENT FILING DATE: 2002-02-01
NUMBER OF SEQ ID NOS: 17
SOFTWARE: Patentin Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 12
US-10-062-257-16
Sequence 16, Application US/10062257
Publication No. US20020128201A1
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity 60.v.
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US-10-062-257-16
                                                                                                                                                                                                                                                                                       ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 HWSYG 5
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3 WSFGI 7
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                                                                                                                                                                                                                                                              TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH
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GENERAL INFORMATION:
APPLICANT: HILDERAND, WILLIAM H.
APPLICANT: HILDERAND, WILLIAM H.
APPLICANT: HILDERAND, WILLIAM H.
APPLICANT: HILDERAND, WILLIAM H.
TITLE OF INVENTION: METHOD AND APPRATUS FOR THE PRODUCTION OF ANTIGENS AND
TITLE OF INVENTION: USES THEREOF
FILE REFERENCE: 6680.034
CURRENT APPLICATION NUMBER: 06/256,410
FRIOR APPLICATION NUMBER: 60/256,410
FRIOR FILING DATE: 2000-12-18
FRIOR FILING DATE: 2000-12-18
FRIOR FILING DATE: 2000-12-18
FRIOR FILING DATE: 2000-12-18
FRIOR FILING DATE: 2001-10-10
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ilarity 75.0%; Pred. No. 5.9e+05;
Conservative 1; Mismatches 0
        PRIOR FILING DATE: 2000-12-12
NUMBER OF SEQ ID NOS: 308
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 72
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; LOCATION: (3)
; OTHER INFORMATION: Unknown amino acid
US-10-022-066-442
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Publication No. US20030059895A1
GENERAL INFORMATION:
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Publication No. US20030166057A1
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APPLICANT: Pia M. Challita-Eid
APPLICANT: Steve Chappell Mitchell
APPLICANT: Daniel B.H. Afar
APPLICANT: Arthur B. Raitano
APPLICANT: Aya Jakobovits
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                                                                                                                                                                                                                                                                                                                       4; Conservative
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; ORGANISM: homo sapien
US-09-791-389-72
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Best Local Similarity
Matches 3; Conserv
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Best Local Similarity
Matches 4; Conserv
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SYGVKP 8
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NAME/KEY: MOD_RES
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                                                                                                                                                                                                                                                                                   sequence 3. Application US/10016283
publication No. US20020164702A1
GENERAL INFORMATION:
APPLICANT: Valenzulate et al., David M.
TITLE OP INVENTION: NOVEL TYROSINE KINASE RECEPTORS AND LIGANDS
FILE REFERENCE: REG195-B-PCT-US
CURRENT APPLICATION NUMBER: US/10/016,283
CURRENT FILING DATE: 2001-11-30
PRIOR APPLICATION NUMBER: US/09/077,955A
PRIOR APPLICATION NUMBER: PCT/US96/20696
PRIOR PPLING DATE: 1998-09-10
PRIOR PILING DATE: 1998-12-13
NUMBER OF SEQ ID NOS: 36
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 3
LENGTH: 6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; Sequence 38, Application US/09990832C
; Publication No. US20030149235Al
; GENERAL INFORMATION:
; APPLICANT: University Court of the University of Glasgow
; TITLE OF INVENTION: Targeting peptides
; FILE REFERENCE: PC/MC/JM/P11910US
; CURRENT APPLICATION NUMBER: US/09/990,832C
; CURRENT FILING DATE: 2003-01-27
; SOFTWARE: Patentin version 3.1
; SEQ ID NO 38
                                                                44.8%; Score 26; DB 15; Length 9; 42.9%; Pred. No. 5.9e+05; Live 2; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           43.1%; Score 25; DB 14; Length 6; 75.0%; Pred. No. 5.9e+05; Live 1; Mismatches 0; Indels
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43.1%; Score 25; DB 12; Length 7;
Best Local Similarity 66.7%; Pred. No. 5.9e+05;
Matches 4; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            , OTHER INFORMATION: Targeting peptide sequence US-09-990-832C-38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: PRT
ORGANISM: Artificial Sequence
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Best Local Similarity 75.0
Matches 3; Conservative
                                                                Query Match
Best Local Similarity 42.5
Matches 3; Conservative
; OTHER INFORMATION: IV. US-10-165-603-8
                                                                                                                                                 1 HWSYGLR 7
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3 WAYG 6
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                                                                                                                                                                                                                                                         RESULT 14
US-10-016-283-3
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Search completed: November 17, 2003, 18:39:40 Job time : 23.5 secs

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Scoring table:

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Result è.

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Sequence 4, Appli
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Sequence 6, Appli
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Sequence 12, Appli
Sequence 12, Appli
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Sequence 7, Appli
Sequence 7, Appli
Sequence 7, Appli
    Sequence 35,
Sequence 36,
Sequence 37,
Sequence 2, #
                                                                                                                                                                                                                                                         Sequence 3
                                                                                                                                                                                                                                              Seguence
                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Zohar, Y.
APPLICANT: Rivier, J.
APPLICANT: Powell, J.
APPLICANT: Sherwood, J.
APPLICANT: Gothilf, Y.
TITLE OF INVENTION: Compounds and Methods For Controlling
TITLE OF INVENTION: Reproduction in Fish
NUMBER OF SEQUENCES: 26
CORRESPONDENCE ADDRESS:
ADDRESSEE: Pennie & Edmonds
STREET: 1155 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 89.7%; Score 52; DB 1; Length 9; Best Local Similarity 88.9%; Pred. No. 2.5e+05; Matches 8; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY: NEW TOIR
COUNTRY: USA
ZID: 10036-22 FORM:
COMPUTER TRADABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Dam PC Compatible
COMPUTER: Patent In Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/341,219
FILING DATE: 05-DEC-1994
CLASSIFICATION NUMBER: 30742
REGISTRATION NUMBER: 30742
REFERENCE/DOCKET NUMBER: 309-003-999
TELEPAN: (212) 790-9090
US-09-450-443E-35
US-09-450-443E-36
US-07-983-111-2
US-07-983-111-4
US-07-983-111-4
US-08-193-374-4
US-08-193-374-4
US-08-193-374-6
US-08-193-374-6
US-08-193-374-6
US-08-193-374-6
US-08-193-374-6
US-08-193-374-9
US-08-332-071B-7
US-08-332-071B-7
US-08-332-071B-7
US-08-332-071B-7
US-08-332-071B-7
US-08-332-071B-7
US-08-332-071B-13
US-08-332-071B-13
US-08-332-071B-13
US-08-332-071B-13
US-08-332-071B-13
US-08-332-071B-13
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TOPOLOGY: unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE: peptide
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      US-08-341-219-8
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US-08-341-219-8
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Sequence 3, Appli
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Sequence 3, Appli
Sequence 1, Appli
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Sequence 1000, Ap
Sequence 1000, Ap
Sequence 4, Appli
Sequence 9, Appli
Sequence 12, Appli
Sequence 12, Appli
Sequence 14, Appli
Sequence 16, Appli
Sequence 20, Appli
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Sequence 2, Appli
Patent No. 5519002
Sequence 10, Appl
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                                                                                                          November 17, 2003, 18:28:20 ; Search time 14.5 Seconds (without alignments) 26.262 Million cell updates/sec
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /cgn2_6/ptodata/1/iaa/5A_COMB.pep:*
/cgn2_6/ptodata/1/iaa/5B_COMB.pep:*
/cgn2_6/ptodata/1/iaa/6A_COMB.pep:*
/cgn2_6/ptodata/1/iaa/RB_COMB.pep:*
/cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*
/cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*
                     GenCore version 5.1.6
(c) 1993 - 2003 Compugen Ltd.
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S-09-082-279B-1000
S-09-315-304B-1000
S-09-834-784-1000
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US-09-450-443E-16
US-09-450-443E-20
US-09-450-443E-22
US-09-450-443E-21
US-09-450-443E-31
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US-08-591-917-2
US-09-309-828-3
US-09-309-828-4
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US-08-871-689-2
US-08-188-918-2
US-08-591-917-3
5488036-3
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US-08-343-883-2
5519002-1
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Maximum Match 100%
Listing first 45 summaries
                                                                                - protein search, using sw model
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Gapop 10.0 , Gapext 0.5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Issued Patents AA:*
                                                                                                                                                                            US-09-462-089-2
58
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Match Length
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Gaps

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ITLE OF INVENTION: peptide, in particular for producing these vaccines...
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                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/343,883
FILING DATE: 17-NOV-1994
CLASSIFICATION 474
FILING DATE: 09-NOV-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/946,495
FILING DATE: 01-MAR-1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: FR 9102513
FILING DATE: 10-DEC-1991
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 8 amino acids

LENGTH: amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE: Luteinizing hormone-relaesing hormone (LH-R) TITLE: activity of some synthetic polypeptides. I. TITLE: Fragments shorter than decapeptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                86.2%; Score 50; DB 1; Lularity 100.0%; Pred. No. 2.5e+05; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   'note= "amidated glycine'
                                                              E: Larson and Taylor
727 Twenty-Third Street, South
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Biochem, Biophys, Res. Commun
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RELEVANT RESIDUES IN SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Schally, A. V. Arimura, A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME/KEY: Peptide
LOCATION: 8
OTHER INFORMATION: /no
OTHER INFORMATION: /no
PUBLICATION INFORMATION: AUTHORS: Schally, A. V
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Yanaihara, N.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AUTHORS: Wissman, H. AUTHORS: Jaeger, G. AUTHORS: Sandow, J. AUTHORS: Yanaihara, N.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 4
5519002-1
; Patent NO. 5519002
; APPLICANT: MIA, ABDUS S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MOLECULE TYPE: peptide
                      NUMBER OF SEQUENCES: CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AUTHORS: Geiger, R. AUTHORS: Konig, W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: amino acid
TOPOLOGY: linear
                                                                                  STREET: 727 Twent CITY: Arlington STATE: Virginia COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            366-375
                                                                ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           JOURNAL:
VOLUME: 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AUTHORS:
AUTHORS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-08-343-883-2
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Matches
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APPLICANT: Roulet, Claude J.M.
APPLICANT: Chouvet, Claire D.
APPLICANT: Bonneau, Michel B.
TYLLE OF INVENTION: Method for improving the organoleptic
TITLE OF INVENTION: qualities of the meat from uncastrated male domestic
TITLE OF INVENTION: animals, vaccines which are usable in this method, new
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ö
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                                                                                                                                                                                                             APPLICANT: Rivier, J.
APPLICANT: Powell, J.
APPLICANT: Sherwood, N.
APPLICANT: Grewood, N.
APPLICANT: Grewood, N.
TITLE OF INVENTION: Compounds and Methods For Controlling
TITLE OF INVENTION: Reproduction in Fish
NUMBER OF SEQUENCES: 26
CORRESPONDENCE ADDRESS:
ADDRESSE: ADDRESSE: Remonds
STREET: 1155 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              89.7%; Score 52; DB 3; Length 9; 88.9%; Pred. No. 2.5e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION DATA:

SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/912,314A

FILING DATE: 30-40N-1997

CLASSIFTCATION: 514

PRIOR APPLICATION DATA:

APPLICATION NUMBER: 08/341,219

FILING DATE: 08-DEC-1994

ATTORNEY/AGENT INFORMATION:

NAME: COTUZZI, Lauta A.

REGISTRATION NUMBER: 8399-003-999

TELECOMMUNICATION NUMBER: 8399-003-999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Pred. No. 2.56
0; Mismatches
                                                                                                                                                                                                                                                                                                                                   JACUSEE: Pennie & Edmonds
STREET: 1155 Avenue of the Americas
CITY: New York
STATE: N.Y.
COUNTRY: 1150
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                            Sequence 8, Application US/08912314A Patent No. 6210927
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; Sequence 2, Application US/08343883
; Patent No. 5573767
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEPHONE: (212) 790-9090
TELEFAX: (212) 869-8864/9741
INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ESS: single
unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity 88.9
Matches 8; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 9 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ZIP: 10036-2711
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy o
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: unknown
MOLECULE TYPE: peptide
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1 HWSYGLSPG 9
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1 HWSYGLSPG 9
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                                                                                  RESULT 2
US-08-912-314A-8
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US-08-343-883-2
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US-09-450-443E-10

US-09-450-443E-10

Sequence 10, Application US/0945043E

Patent No. 6566402

GENERAL INFORMATION:

APPLICANT: DELANSORNE, Remi

APPLICANT: DELANSORNE, Remi

TITLE OF INVENTION: LH-RH peptide analogues, their uses and pharmaceutical

TITLE OF INVENTION: LW-RH peptide analogues, their uses and pharmaceutical

TITLE OF INVENTION: LW-RH peptide analogues, their uses and pharmaceutical

TITLE OF INVENTION: LW-RH peptide analogues, their uses and pharmaceutical

FILE REFERENCE: 12058-405

CURRENT APPLICATION NUMBER: US/09/450,443E

CURRENT APPLICATION NUMBER: PCT/EP98/02802

PRIOR FILING DATE: 1998-05-06

PRIOR FILING DATE: 1997-06-02

NUMBER OF SEQ ID NOS: 39

SOFTWARE: PatentIN Ver. 2.1

SEQ ID NO 10

LENGTH: 9
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OTHER INFORMATION: Description of Artificial Sequence: LH-RH analogue
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0. 2.5e+05; o; Indels
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Best Local Similarity 87.5%; Pred. No. 2.5e+05;
Matches 7; Conservative 0; Mismatches 1;
TITLE OF INVENTION: METHOD AND COMPOSITION FOR
                                                                                                                                                                                                                                                                                              86.2%; Scc...
100.0%; Pred. No. ...
0; Mismatches
               ; PREVENTING CONCEPTION
; CURRENT APPLICATION DATA:
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/26,180
FILING DATE: 01-MAR-1993
; PRIOR APPLICATION NUMBER: 697,127
; FILING DATE: 08-MAY-1991
; FILING DATE: 29-SEP-1988
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 2, Application US/08591917
Patent No. 5707964
GENERAL INFORMATION:
APPLICANT: Nett, Torrance M
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OTHER INFORMATION: Xaa is pGlu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: PRT ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; LOCATION: (7)
; OTHER INFORMATION: Xaa is Npg
US-09-450-443E-10
                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity 100.
Matches 8; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                     2 WSYGLRPG 9
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NAME/KEY: SITE
LOCATION: (7)
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                                                                                                                                                                                                                             SEQ ID NO:1:
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Sequence 3, Application US/09309828
; Sequence 3, Application US/09309828
; Sequence 3, Application US/09309828
; GENERAL INPORMATION:
    APPLICANT: Jensen, Knud J.
    APPLICANT: Barany, George
; APPLICANT: Songster, Micheal F.
    APPLICANT: Albaricio, Fernando
; APPLICANT: Albaricio, Fernando
; APPLICANT: Vagner, Josef
; TITLE OF INVENTION: SUPPORT MATERIAL FOR SOLID PHASE ORGANIC SYNTHESIS
; TITLE OF INVENTION: SUPPORT MATERIAL FOR SOLID PHASE ORGANIC SYNTHESIS
; TITLE OF INVENTION: SUPPORT WATERIAL FOR SOLID
; FILE REFERENCE: 110.00220102
; CURRENT APPLICATION NUMBER: US/09/309,828
; CURRENT PILING DATE: 1999-05-11
; SEQ ID NOS: 4
; SOFTWARE: Patentin Ver. 2.1
; SEQ ID NO 3
; LENGTH: 9
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OTHER INFORMATION: Description of Artificial Sequence:Des-Glylo,
OTHER INFORMATION: methylamide9-LHRH
NAME/KEY: SITE
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OMPRATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/591,917
FILING DATE: 26-JAN-1996
CLASSIFICATION: 514
APPLICANT: Glode, Leonard Michael
TITLE OF INVENTION: A METHOD FOR TREATING CANCER
NUMBER OF SEQUENCES: A
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 45; DB 1; I
Pred. No. 2.5e+05;
0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ATTORNEY AGENT INPORMATION:
NAME: Kovarik, Joseph E.
REGISTRATION NUMBER: 33,005
REFERENCEJOCKET NUMBER: 2730-3-2-1-1
TELECOMMUNICATION INPORMATION:
TELEPHONE: (303) 863-023
INFORMATION FOR SEQ ID NO: 2:
INFORMATION FOR SEQ ID NO: 2:
LENGTH: 9 amino acids
                                                                                          ADDRESSEE: Sheridan Ross & McIntosh
STREET: 1700 Lincoln Street, Suite 3500
CITY: Denver
STATE: Colorado
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: PRT
ORGANISM: Artificial Sequence
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Best Local Similarity 87.5
Matches 7; Conservative
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MOLECULE TYPE: protein

US-08-591-917-2
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OTHER INFORMATION: pGlu
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Gaps
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US-08-871-689-2
Sequence 2, Application US/08871689
Sequence 2, Application US/08871689
Sequence 2, Application US/08871689
September 1000 Septemb
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Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SIGNE: VIKULINIA
CUCUNIKY: VIKULINIA
Z19: 22314
COMPUTER READABLE FORM:
MEDIUW TYBE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/871,689
FILING DATE: 09-JUN-1997
CLASSIFICATION: 424
PRIOR APPLICATION ADATA:
APPLICATION NUMBER: US 08/185,878
ATTORNEY/AGRYT INFORMATION:
NAME: MILLS, DEMETRA J.
REGISTRATION NUMBER: 34,506
REGISTRATION NUMBER: 34,506
REGISTRATION NUMBER: 34,506
REGISTRATION NUMBER: 34,506
REGISTRATION NUMBER: 1451-004
TELECOMMUNICATION INFORMATION:
TELEFRAK: 703-684-1111
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100.0%; Pred. No. 2.5e+05;
ative 0; Mismatches 0;
Mismatches
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Patent No. 5480656
GENERAL INPRMATION:
APPLICANT: OKADA, Hiroaki
APPLICANT: INOUE, Yayoi
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Best Local Similarity 100.
Matches 7; Conservative
7; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE: protein HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   amino acid
                                                             1 HWSYGLRP 8
                                                                                                                          2 HWSYXLRP 9
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   Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Jensen, Knud J.
APPLICANT: Barany, George
APPLICANT: Barany, George
APPLICANT: Songster, Micheal F.
APPLICANT: Albaria, Jordi
APPLICANT: Alsina, Jordi
APPLICANT: Vagner, Josef
TITLE OF INVENTION: SUPPORT MATERIAL FOR SOLID PHASE ORGANIC SYNTHESIS
FILE REFERENCE: 110.00220102
CURRENT APPLICATION NUMBER: US/09/309,828
CURRENT FILING DATE: 1999-05-11
                                                                                                                          Gaps
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; Patent No. 5488036
; Patent No. 5488036
; TITLE OF INVENTION: METHOD FOR STERLLIZING ANIMALS USING HORMONE-TOXIN CONUGATE COMPOUNDS
; NUMBER OF SEQUENCES: 5
; NUMBER OF SEQUENCES: 5
; CURRENT APPLICATION DATA: 4PPLICATION NUMBER: US/08/94,625
FILING DATE: 20-70L-1993
; PRIOR APPLICATION NUMBER: 837,639
FILING DATE: 14-FEB-1992
; FILING DATE: 23-FEB-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           77.6%; Score 45; DB 4; Length 9; 100.0%; Pred. No. 2.5e+05; cive 0; Mismatches 0; Indels
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                                                         77.6%; Score 45; DB 4; Length 9; 100.0%; Pred. No. 2.5e+05; cive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                              Sequence 4, Application US/09309828
Patent No. 6566494
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87.58;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; OTHER INFORMATION: Pro-NHEC US-09-309-828-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQ ID NOS: 4
SOFTWARE: PatentIn Ver. 2.1
                                                             Query Match 77.6
Best Local Similarity 100.
Matches 7; Conservative
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Best Local Similarity 100.
Matches 7; Conservative
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Best Local Similarity
                                                                                                                                                                                       1 HWSYGLR 7
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2 HWSYGLR 8
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LOCATION: (1)
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US-09-309-828-4
US-09-309-828-3
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LENGTH: 9
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Gaps
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PAPLICANT: NETT, TORRANCE M.;GLODE, LEONARD M.

TITLE OF INVESTION: METHOD FOR STERILIZING ANIMALS USING HORMONE-TOXIN CONJUGETE COMPOUNDS

NUMBER OF SEQUENCES: 5

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/94,625

FILING DATE: 20-JUL-1993

PRIOR APPLICATION DATA:

APPLICATION NUMBER: 837,639

FILING DATE: 14-FEB-1992

PRING DATE: 23-FEB-1992
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Fatent No. 5480656
GENERAL INFORMATION:
APPLICANT: OCADA, Hiroaki
APPLICANT: OCADA, Yasuaki
INGUENT: OCAMA, Yasuaki
ITILE OF INVENTION:
CORRESPONDENCES: 2
CORRESPONDENCE ADDRESS:
ADDRESSE: WEGNER, CANTOR, WUELLER & PLAYER
STREET: 1233 20th Street, N.W., Suite 300
CITY: Washington
STREET: 0.C.
COUNTRY: U.S.A.
ZIP: 20036-8218
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Pred. No. 2.5e+05;
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Pred. No. 2.5e+05;
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MEDIUM TYPE: Floppy disk
COMPUTER: COMPAQ 286 IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                          NAME: KOVARIK, JOSEPH E.
REGISTRATION NUMBER: 33,005
REFRENCE/DOCKET NUMBER: 2730.
TELEPHONE: (303) 863-9700
TELEPHONE: (303) 863-9700
TELEPHONE: (303) 863-023
INVERMATION FOR SEQ ID NO: 3: SEQUENCE CHARACTERISTICS:
        ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity 87.5
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Best Local Similarity 87.5
Matches 7; Conservative
                                                                                                                                                                                                                                                                                        TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
US-08-591-917-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 HWSYGLRP 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2 HWSYKLRP 9
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US-08-188-918-1
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APPLICANT: OGAWA, Yasuaki
TITLE OF INVENTION: PROLONGED RELEASE MICROCAPSULES
NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
ADDRESSEE: WEGNER, CANTOR, MUELLER & PLAYER
STREET: 1233 20th Street, N.W., Suite 300
CITY: Washington
STRATE: D.C.
COUNTRY: U.S.A.
ZIP: 20036-8218
COMPUTER: COMPAD 286 IBM PC compatible
COMPUTER: D.C.
COUNTRY: U.S.A.
ZIP: 20036-8218
COMPUTER: D.C.
COUNTRY: U.S.A.
SOFTWARE: Patentin Release #1.0, Version #1.25
COURRENT APPLICATION NUMBER: US/08/188,918
FILING DATE: 31-JAN 1994
CLASSIFICATION NUMBER: US/08/188,918
FILING DATE: 02-FEB-1990
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 9;
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APPLICANT: Nett, Torrance M
APPLICANT: Glode, Leonard Michael
TITLE OF INVENTION: A METHOD FOR TREATING CANCER
NUMBER OF SEQUENCES: 3
CORRESPONDENCE ADDRESS:
ADDRESSEE: Sheridan Ross & McIntosh
STREET: 1700 Lincoln Street, Suite 3500
CITY: Denver
STATE: Colorado
COUNTRY: USA
ZIP: 80203
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/591,917
FILING DATE: 26-JAN-1996
CLASSIFICATION: 514
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 44; DB 1; I
Pred. No. 2.5e+05;
0; Mismatches 1;
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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Patent No. 5707964
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Best Local Similarity 97.5
Matches 7; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE: peptide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           amino acid
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Search completed: November 17, 2003, 18:32:38 Job time: 15 secs

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             KESULT 15
US-09-082-279B-1000
SGRUECE 1000, Application US/09082279B
Fatent No. 6258782
Fatent No. 6258782
GENERAL INFORMATION:
APPLICANT: Barney, Shawn
APPLICANT: Guthris, Kelly
APPLICANT: Anwer, Mohmed
APPLICANT: Lambert, Dennis
TITLE OF INVENTION: HYBRID POLYPEPTIDES WITH ENHANCED
TITLE OF INVENTION: HYBRID POLYPEPTIDES
TILE REFERENCE: 7872-043
CURRENT APPLICANT: 1998-05-20
NUMBER OF SEQ ID NOS: 1515
SOFTWARE: FestSEQ for Windows Version 3.0
SEQ ID NO 1000
LEMETHER OF SELECTION WINDOWS: 1515
SEQ ID NO 1000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME/KEY: Modified-site
LOCATION: 6
COTHER INFORMATION: /label= D-aminoacid
OTHER INFORMATION: /note= "D-aminoacid at position 6"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     72.4%; Score 42; DB 1; Length 9; 87.5%; Pred. No. 2.5e+05; Live 0; Mismatches 1; Indels
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/188,918
FILING DATE: 31-JAN-1994
CLASSIFICATION: 424
PRIOR APPLICATION NUMBER: US 033133-1990
FILING DATE: 02-FEB-1990
ATTORNEY/AGENT INFORMATION:
NAME: MUELLER, DOUG148 P.
REGISTRATION NUMBER: 20,300
REFERENCE/DOCKET NUMBER: P8700-22697A
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEFAX: (202) 885-0605
ITELEX: 440706 WEGBR
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LUNGTH: 9 amino acids
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: peptide
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity 87.5
Matches 7; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 HWSYGLRPG 9
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2 HWSYLLRP 9
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1 HWSY-LRPG 8

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pseudopeptide

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Luteinizing hormon Gonadotrophin rele Luteinizing hormon Viral core polypep Luteinizing hormon Leader sequence 46

Luteinising hormon LH and FSH releasi LH and FSH releasi

OM protein

Run on:

Sequence:

Searched:

Database

Uhmann R, Radscheit K;

24406786

Result Š.

Self adjuvanating
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Alpha-substituted
Alpha-substituted
Alpha-substituted
Cholecystokinin an
Cholecystokinin an
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Examples; Page 30; 41pp; English.
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                                                                                                 8; Conservative
                                                                                                                                                                                                                                                      LHRH peptide fragment.
                                                                                                                     8
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                                                                              Query Match
Best Local Similarity
                                                                                                                                         1 WSYGLRPG
                                                                                                                     WSYGLRPG
                                                          8 AA;
                                                                                                                                                                                                                                  11-MAY-1999
                                                                                                                                                                                                                                                                                                                                                         WO9902180-A1
                                                                                                                                                                                                                                                                                                                                                                                               09-JUL-1998;
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                                                                                                                                                                                                                                                                                                                                     Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                           McNamara MK;
                                                                                                                                                                                                                                                                                                                                                                             21-JAN-1999.
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                                                                                                                                                                                                              AAW94892;
                                                          Sequence
                                                                                                 Matches
                                                                                                                                                                       RESULT 3
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                                                                              of
                                                                                                                                                                       Gaps
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                                                                                                                                                                                                                                                                                                                                                        Immunoneutralisation; luteinising hormone releasing hormone; GnRH;
                                                                             The peptides of the invention are intermediates for the synthesis gonadorelin (luteinising hormone releasing hormone) and its analogues (see e.g. US 4024248). (Updated on 16-AUG-2002 to add missing OS field.)
                            Prodn. of gonadorelin peptide intermediates without racemisation - from new protected tryptophan tri:peptide derivs.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Improving meat quality of intact male animals - by immuno:neutralisation, shortly before slaughter, of steroid with anti-LHRH, esp. induced by two-stage vaccination
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                                                                                                                                                    Length 8;
                                                                                                                                                                      Indels
                                                                                                                                                100.0%; Score 50; DB 6; L. 100.0%; Pred. No. 9.3e+05; rive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Roulet C;
                                                                                                                                                                                                                                                                                                                                                                                                         Location/Qualifiers
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                                                                                                                                                                                                                                                                AAR26733 standard; peptide; 8 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 22; Page 17; 18pp; French
                                                          Claim 4; Page 23; 28pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        92EP-0400496.
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(first entry)
                                                                                                                                                                                                                                                                                                                                                                  gonadoliberin; castration
                                                                                                                                                 Query Match 100.
Best Local Similarity 100.
Matches 8; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Chouvet C,
                                                                                                                                                                                                                                                                                                                                   Immunogenic LHRH(3-10).
         WPI; 1985-243923/40.
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                                                                                                                                                                                                        WSYGLRPG 8
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                                                                                                                                8 AA;
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Modified-site
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11-FEB-1993
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                                                                                                                                                                                                                                                                                                                                                                                      Synthetic.
                                                                                                                                                                                                                                                                                    AAR26733;
                                                                                                                                 Sequence
                                                                                                                                                                                                                                           RESULT 2
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The invention relates immunogenic composition for eliciting an immune response to luteinising hormone releasing hormone (LHRH). The composition comprises a LHRH-diphtheria toxoid (DT) conjugate adsorbed to an ionic polygaccharide. The LHRH-DT compositions can be used for eliciting an immune response to LHRH, for castrating an animal, for regulating osertus cycling in a female animal or for inhibiting characteristics induced by the sexual maturation of an animal, e.g. aggression or sexual activity. They can also be used for achieving corpanolophic characteristics from the meat of livestock. They can also be used for inhibiting the growth of cells which are regulated directly or used for inhibiting the growth of cells which are regulated directly or indirectly by LHRH, e.g. malignant breast cells, malignant covarian cells, malignant concocetal cells or hyperplastic cells. They can also be used for down-regulating the libido of an animal. They can also be used for inhibiting pregnancy, prostate cells animal. They can also be used for inhibiting pregnancy, prostate cells animal. They can also be used for inhibiting pregnancy, prostate cells animal andometriosis or inhibiting pregonses. The LHRH compositions induce a more effective immune response against LHRH than
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leaber carcasses) to be retained practically up to the time of slaughter. The treatment does not induce any local reactions which could result in the meat being rejected on grounds of quality. (Updated on 25-MAR-2003 to correct PN field.) (Updated on 25-MAR-2003 to correct PA field.)
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WO9302706-A1
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15-JUN-1993
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Finney KG,
                                                                                                                                                                                                                                                                                                              Synthetic.
                                                                                                                                 AAR32891;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present sequence is a peptide fragment of human luteinising hormone releasing hormone (also known as LHRH, GnRH and gonadotrophin releasing hormone). It was used to demonstrate the novel adjuvant of the invention, which has lower reactogenicity then previous compositions. Vaccination of humans and animals against LHRH can be used as a method of fertility control, as well as enabling the control and treatment of disorders of the reproductive organs, such as testicular, breast, prostate and ovarian cancers, prostate enlargement and endometriosis. The composition of the larter of which is an immunostimulant, and it can also be used with other immunogens including soluble protein antigens, peptide haptens conjugated to a carrier protein and whole viruses.
                    the hormones LH and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Adjuvant composition for manufacturing an immunogenic composition that can elicit an immune response in animmal, comprises an ionic polysaccharide component and a saponin component that is an immunostimulating complex.
   The effective immune response
                                                                                                                                                  Gaps
                 against LHRH results in prevention of the release of the hormones 1
FSH from the anterior pituitary. Sequences AAW94890-93 are peptide
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                                                                                                                Length 8;
                                                                                                                                              Indels
                                                                                                              100.0%; Score 50; DB 20;
100.0%; Pred. No. 9.3e+05;
tive 0; Mismatches 0;
   the LHRH-carrier-adjuvant compositions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure; Page 50; S3pp; English.
                                                                                                                                                                                                                                                                                               AAB15364 standard; peptide; 8 AA.
                                                                                                                                                                                                                                                                                                                                                                                              Human LHRH peptide SEQ ID NO: 3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               99WO-AU01167.
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                                                                                                                                                  Conservative
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                                                 derivatives of LHRH
                                                                                                              Query Match
Best Local Similarity
Matches 8; Conserv
                                                                                                                                                                                 1 WSYGLRPG
                                                                                                                                                                                                             1 WSYGLRPG
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                                                                                 8 AA;
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Gonadotrophin releasing hormone agonist; GnRH agonist; combination drug;
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                                                                                                                                                                                                                       Luteinising hormone releasing hormone; chemical sterilisation; antibody response; AIDS; malaria; cancer; breast; prostate; endometrial; ovarian.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Self-adjuvanating vaccines - comprises peptide conjugated to fatty acids and carrier, useful for treating prostate, breast, endometrial and ovarian cancers, etc.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Jennings PA, Reilly WG, Whittaker RG;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     e.g. AIDS, malaria, etc. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                    Self adjuvanating vaccine LHRH peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (CSIR ) COMMONWEALTH SCI & IND RES ORG.
AAR32891 standard; peptide; 7 AA.
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                                                                                                                                    (first entry)
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Best Local Similarity 100.
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Local Similarity 100.0

Query Match Matches

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Gaps

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Indels

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100.0%; Score 50; DB 21; Length 8; 100.0%; Pred. No. 9.3e+05;

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Self adjuvanating vaccine LHRH peptide
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                                                                                                                                                                                                                                             18-FEB-1993.
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                                                                                                                                                                                                                                                                                                                                                                                                                                     Finney KG,
                                                                                                                                              Synthetic.
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Best Local S
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THE STATE OF THE S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention describes a pharmaceutical composition (1) for treating breast cancer, endometriosis, myometrium tumour, Alzheimer's disease, circulatory system disorders, irregular disease, circulatory system disorders, irregular disease, circulatory system disorders, irregular periods cancer metastasis, premenstrual syndrome, muscular distress or osteopathies due to calcium/phosphorus imbalance. (1) comprises a confectrophin releasing hormone (GRRM) agonist and a selective osetrogen receptor modulator (SERM). (1) has cytostatic, gynaecological, nootropic, neuroprotective, cardiovascular, endocrine, muscular and osteopathic activities. (1) can be used as GRRH agonists used in combination with selective osetrogen receptor modulators, selective androgen receptor modulators, selective androgen receptor modulators, lyase inhibitors, receptor-type tyrosine kinase inhibitors, lyase inhibitors, receptor entegonise kinase inhibitors, lyase inhibitors, receptor treating and preventing endomertiosis myometrium tumour, antagonists for treating and preventing endomertiosis, myometrium tumour, are disease, circulatory system disorders, menopausal disorders, irregular periods, cancer metastasis, premenstrual syndrome, muscular distress, osteopathies due to calcium/phosphorus imbalance, prostate cancer cor breast cancer or their reoccurrence or metastasis. Combination enhances the quality of life by enhancing cor metastasis. Combination enhances the quality of life by enhancing correspondent agonist related peptide which is given in the present
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                      Alzheimer's disease, circulatory system disorder; menopausal disorder; irregular period; cancer metastasis; premenstrual syndrome; osteopathy; muscular distress; calcium/phosphorus imbalance; SERM; gynaecological; selective oestrogen receptor modulator; cytostatic; nootropic; muscular; neuroprotective; cardiovascular; endocrine; osteopathic; prostatomegally;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Pharmaceutical composition e.g. for breast cancer comprises gonadotrophin releasing hormone agonist and selective estrogen receptor
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    pharmaceutical; breast cancer; endometriosis; myometrium tumour;
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                                                                                                                                                                                                                                                                                                                                           08-AUG-2002; 2002WO-JP08130
                                                                                                                                                                                                                                                                                                                                                                                        10-AUG-2001; 2001JP-0244616
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(first entry)
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Best Local Similarity
7; Conserve
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WSYGLRP 8
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                                                                                                                                                   prostate cancer.
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15-JUN-1993
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                                                                                                                                                                                             Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Furuya S,
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Luteinising hormone releasing hormone (LH-RH) related peptide SEQ ID:156.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The sequence is that of a luteinising hormone releasing hormone (LHRH) peptide which can be used as part of a self adjuvanating vaccine for raising antibodies to the LHRH peptide. The vaccine may be used in the chemical sterilisation of animals and for treating breast, prosester, endometrial and ovarian cancer. Using peptide epitopes of AIDS, malaria, influenza, hepatitis or zona pellucida the vaccine can be used to enhance the immune response of an animal to the antigen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Protection, endogenous therapeutic peptide, peptidase, conjugation, blood component, modification, succinimidyl; maleimido group; amino, hydroxyl; thiol; hormone; growth factor; neurotransmitter.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ;
Luteinising hormone releasing hormone; chemical sterilisation; antibody response; AIDS; malaria; cancer; breast; prostate;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Self-adjuvanating vaccines - comprises peptide conjugated to fatty acids and carrier, useful for treating prostate, breast, endometrial and ovarian cancers, etc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 40; DB 14; Lengtn ,,
Pred. No. 9.3e+05;
Pred. No. 9.3e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Reilly WG, Whittaker RG;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            e.g. AIDS, malaria, etc. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (CSIR ) COMMONWEALTH SCI & IND RES ORG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAB90982 standard; Peptide; 7 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 7; Page 15; 37pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            80.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                 91AU-0007457.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             17-MAY-2000; 2000WO-US13576
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               99US-0134406
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Jennings PA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               6; Conservative
                                                                         endometrial; ovarian.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1993-076183/09.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 WSYGLQP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 WSYGLRP
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Luteinising hormone releasing hormone (LH-RH) related peptide SEQ ID:150.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Modifying and attaching therapeutic peptides to albumin prevents peptidase degradation, useful for increasing length of in vivo activity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Protection, endogenous therapeutic peptide, peptidase, conjugation, blood component, modification, succinimidyl, maleimido group, amino, hydroxyl, thiol; hormone, growth factor, neurotransmitter.
                                                                                                                                                       Prodn. of gonadorelin peptide intermediates without racemisation - from new protected tryptophan tri:peptide derivs.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Thibaudeau K;
                                                                                                                                                                                                                                                      the
                                                                                                                                                                                                                                                                                                                                                                                                                              1; Indels
                                                                                                                                                                                                                                                                                                                                                                                          Length 7;
                                                                                                                                                                                                                                                  The peptides of the invention are intermediates for gonadorelin (luteinising hormone releasing hormone) analogues (see e.g. US 4024248). (Updated on 16-AUG-2002 to add missing OS field.)
                                                                                                                                                                                                                                                                                                                                                                                        Score 38; DB 6; I
Pred. No. 9.3e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Holmes DL,
                                                                                                                                                                                                                                                                                                                                                                                                                                0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Page 239; 733pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ezrin AM, Milner PG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAB90976 standard; Peptide; 8 AA.
                                                                                                                                                                                                                Claim 5; Page 23; 28pp; German.
                                                                                                                                                                                                                                                                                                                                                                                      76.0%;
85.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    99US-0134406.
99US-0153406.
99US-0159783.
  84DE-3411224.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       22-JUN-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                6; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CONJ-) CONJUCHEM INC
                                                                            Uhmann R, Radscheit
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2001-112059/12.
                                                                                                                 WPI; 1985-243923/40
                                      (FARH ) HOECHST AG.
                                                                                                                                                                                                                                                                                                                                                                                                          Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 WSYSLRP 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 WSYGLRP
                                                                                                                                                                                                                                                                                                                                                   7 AA;
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  27-MAR-1984;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        17-MAY-1999;
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15-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           23-NOV-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic
                                                                                                                                                                                                                                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAB90976;
                                                                                                                                                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 10
AAB90976
                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           임
                                                                                                                                                                                                                                                                                    The present invention describes a modified therapeutic peptide (I) comprising a therapeutically active amino acid region (III) and a reactive group (II) (e.g. succinimidyl and maleimido groups) attached to a less therapeutically active amino acid region (IV), which covalently bonds with amino/hydroxyl/thiol groups on blood components to form a peptidase stabilised therapeutic peptide composed of 3-50 amino acids. (I) are useful for modifying therapeutic peptides e.g. hormones, growth a cactors and neurotransmitters, to protect them from peptidase activity in vivo for the treatment of various disorders. Endogenous therapeutic peptides are not suitable as drug candidates as they require frequent administration due to rapid degradation by peptidases in the body. Modifying and attaching therapeutic peptides to albumin prevents or reduces the action of peptidases to increase length of activity (half life) and specificity as bonding to large molecules decreases. ABB90829 to ABB92411 represent invention.

Company of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ö
                                                                                                                                                                       Modifying and attaching therapeutic peptides to albumin prevents peptidase degradation, useful for increasing length of in vivo activity
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0
                                                                                            Thibaudeau K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gonadorelin; hormone; luteinising hormone releasing hormone.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              note= "bonded to urethane-protecting gp."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   78.0%; Score 39; DB 22; Length 7; 100.0%; Pred. No. 9.3e+05; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
                                                                                               Holmes DL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence of gonadorelin peptide intermediate.
                                                                                                                                                                                                                                                      Disclosure; Page 241; 733pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /label= Pro-NHC2H5
                                                                                            Milner PG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /label= D-Ser(But)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAP50693 standard; peptide; 7 AA.
99US-0153406
99US-0159783
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(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity 100
                                                        (CONJ-) CONJUCHEM INC
                                                                                               Bridon DP, Ezrin AM,
                                                                                                                                   WPI; 2001-112059/12.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2 SYGLRPG 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              7 AA;
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  10-SEP-1999;
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                  15-OCT-1999;
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16-OCT-1991
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Sequence

Query Match Best Loc Matches

02-OCT-1985 EP156280-A

Synthetic

AAP50693;

RESULT 9 AAP50693

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Gaps

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24-MAY-1991
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07-AUG-1989;
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                                                                                                                                                                                                                                                                                        antagonist
                                                                                                                                                                                                                                                                                                            Synthetic.
                                                                                                                                                                                                              AAR11092;
                                                       Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
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                                                                                                                                                                      RESULT 12
                                                                                                                                                                              AAR1109;
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peptidase stabilised therapeutic peptide composed of 3-50 amino acids.

(I) are useful for modifying therapeutic peptides e.g. hormones, growth factors and neurotransmitters, to protect them from peptidase activity in vivo for the treatment of various disorders. Endogenous therapeutic peptides are not suitable as drug candidates as they require frequent administration due to rapid degradation by peptidases in the body. Modifying and attaching therapeutic peptides to albumin prevents or reduces the action of peptidases to increase length of activity (half life) and specificity as bonding to large molecules decreases intracellular uptake and inteference with physiological processes. AAB90829 to AAB2241 represent peptides which can be used in the exemplification of the present invention.
                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This is one of 57 specific examples of highly generic reduced size LHRH analogues. It is based on amino acid residues 3 to 9 of natural LHRH. The analogues can be used for treating diseases
                                                                                                                                                                                ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Reduced size Pseudo peptide LHRH analogues - used as LHRH agonists or antagonists eg in treatment of prostate cancer and benign prostatic hypertrophy
                                                                                                                                                                                                                                                                                                                                                               Leutinizing hormone releasing hormone; pseudopeptide; agonist;
                                                                                                                                                                                                                                                                                                                                                                                                                                    label= N-3-(4-fluorophenyl)propionyl-D-Trp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             label= N-3-(4-fluorophenyl)propionyl-D-Trp
                                                                                                                                                           Score 38; DB 22; Length 8; Pred. No. 9.3e+05; 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Fitzpatrick TD;
                                                                                                                                                                                                                                                                                                                                                                                                                Location/Qualifiers
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                                                                                                                                                                                                                                                                            AAR11091 standard; Protein; 7 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                          label= N-Me-Ser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /label= Pro-NHEt
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 4; Page 87; 90pp; English.
                                                                                                                                                                                                                                                                                                                                          LHRH pseudopeptide analogue #3
                                                                                                                                                         76.0%;
85.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                90EP-0114752,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    90US-0548511.
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                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                           Query Match 76.0
Best Local Similarity 85.7
Matches 6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Palabrica CA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 1991-081768/12.
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                                                                                                                                                                                                                         WSYSLRP 8
                                                                                                                                                                                                    WSYGLRP
                                                                                                                                        8 AA;
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07-AUG-1989;
                                                                                                                                                                                                                                                                                                                      24-MAY-1991
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                                                                                                                                                                                                                                                                                                                                                                          antagonist
                                                                                                                                                                                                                                                                                                                                                                                             Synthetic.
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                                                                                                                                                                                                                                                                                                 AAR11091;
                                                                                                                                         Sequence
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                                                                                                                                                                                                                                                       RESULT 11
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resulting from gonadal hormone over- or under-production in either sex, controlling reproduction, as fertility promoters when administered in pulses or for reducing dihydrotestosterone levels. See also AAR11089-R11090 and AAR11092-4.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Leutinizing hormone releasing hormone; pseudopeptide; agonist;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Reduced size Pseudo peptide LHRH analogues - used as LHRH agonists or antagonists eg in treatment of prostate cancer and benign prostatic hypertrophy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           'label= N-3-(4-fluorophenyl)propionyl-D-Trp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  72.0%; Score 36; DB 12; Length 7;
                                                                                                                                                                                                                                         Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Fitzpatrick TD;
                                                                                                                                                                                     Score 36; DB 12; L
Pred. No. 9.3e+05;
0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                'label= D-Lys(Nic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAR11092 standard; Protein; 7 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               'label= N-Me-Tyr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /label= Pro-NHEt
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LHRH pseudopeptide analogue #4.
                                                                                                                                                                                        72.0%;
85.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        90EP-0114752
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89US-0390269
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                                                                                                                                                                                                                                            6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Haviv F, Palabrica CA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1991-081768/12.
                                                                                                                                                                                        Query Match
Best Local Similarity
Matches 6; Conserv
                                                                                                                                                                                                                                                                                            1 WSYGLRP
                                                                                                                                                                                                                                                                                                                                             WSYWLRP
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                                                                                                                                       7 AA;
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Modified-site
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polymer;

Luteinizing hormone-releasing hormone; LH-RH; cytostatic; gynaecologizal; sustained release microsphere; biodegradable polymer hormone dependent disease; pancreatic cancer; prostatic hypertrophy; breast cancer; endometriosis; myome of the uterus;

neurogenic precocious puberty; contraceptive; deslorelin

Location/Qualifiers

Luteinizing hormone-releasing hormone (LH-RH) agonist, deslorelin.

(first entry)

21-MAY-2002

His-CO-2-pyrrolidone (The CO group is to the 2-pyrrolidone at carbon 5"

"OTHER=

/label= OTHER

Key Modified-site

Synthetic

/note= "OTHER= Pro-NH-CH2-CH2-NH2'

OTHER

8 /label= (/note=

Modified-site

EP1142567-A2 10-OCT-2001 Nagai A;

Ohtani S,

Takechi N,

WPI; 2002-091561/13.

95JP-0327690 96EP-0309136

15-DEC-1995; 13-DEC-1996;

13-DEC-1996; 2001EP-0116787

(TAKE) TAKEDA CHEM IND LTD

AAU76988 standard; Peptide; 8 AA.

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Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         This is one of 57 specific examples of highly generic reduced size LHRH analogues. It is based on amino acid residues 2 to 9 of natural LHRH. The analogues can be used for treating diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            resulting from gonadal hormone over- or under-production in either sex, controlling reproduction, as fertility promoters when administered in pulses or for reducing dihydrotestosterone levels. See also AAR11099 and AAR11091-4.
             .
0
                                                                                                                                                                                                                     Leutinizing hormone releasing hormone; pseudopeptide; agonist;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                agonists or antagonists eg in treatment of prostate cancer and
benign prostatic hypertrophy
                                                                                                                                                                                                                                                                                                          'label= (N-(N-(alpha)-morpholinocarbonyl)-Phe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Reduced size Pseudo peptide LHRH analogues - used as LHRH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 8;
             Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Greer J, Fitzpatrick TD,
 Pred. No. 9.3e+05;
0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 36; DB 12; Le
Pred. No. 9.3e+05;
0; Mismatches 1;
                                                                                                                                                                                                                                                                                  Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                         /label= Pro-NH-Et
                                                                                                                        AAR11090 standard; Protein; 8 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 4; Page 86; 90pp; English.
                                                                                                                                                                                               LHRH pseudopeptide analogue #2
                                                                                                                                                                                                                                                                                                                                  label= D-Trp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   72.0%;
85.7%;
85.78;
                                                                                                                                                                                                                                                                                                                                                                                                                                 90EP-0114752.
                                                                                                                                                                                                                                                                                                                                                                                                                                                        90US-0548511
89US-0390269
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                                                                                                                                                                        (first entry)
             6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Palabrica CA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1991-081768/12.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
Matches 6; Conserv
                                   1 WSYGLRP 7
 Best Local Similarity
                                                         WSYKLRP 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           8 AA;
                                                                                                                                                                                                                                                                                             Modified-site
                                                                                                                                                                                                                                                                                                                     Modified-site
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                                                                                                                                                                                                                                                                                                                                                                                                                                 01-AUG-1990;
                                                                                                                                                                       24-MAY-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                        10-JUL-1990;
07-AUG-1989;
                                                                                                                                                                                                                                                                                                                                                                                                        20-MAR-1991
                                                                                                                                                                                                                                                                                                                                                                                 EP417454-A.
                                                                                                                                                                                                                                    antagonist
                                                                                                                                                                                                                                                          Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Haviv F,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence
                                                                                                                                                AAR11090;
            Matches
                                                                                                           AAR11090
ID AAR1
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The intention describes a subtained release intropher invention describes as substance and a biodegradable polymer, obtained by subjecting a water/oil/water or oil/water emulsion to an in-water drying under particular conditions. The physiologically active substance is in an inner aqueous phase and the biodegradable polymer is in an external oil phase. The sustained release microsphere is useful as a preparation for treatment or prophylaxis of hormone dependent diseases such as pancreatic cancer, prostatic hypertrophy, breast cancer, a contraceptive. The method increases the rate of solvent removal from microspheres, reducing the amount of solvent in microspheres in a short time and markedly improves the drug entrapment ratio in microspheres by subjecting the microspheres are excellent in toxicity and can be used safely. Microspheres are excellent in contactity at the time of collection and in dispersibility and needle passability when they are used as a medicinal injectable preparation. This sequence represents a peptide agonist of the luteinizing contacts the contact of the creation of microspheres.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Production of a sustained-release microsphere having physiologically active substance and a biodegradable polymer involves subjecting a water/oil/water or oil/water emulsion to an in-water drying under particular conditions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention describes a sustained release microsphere having a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; Page 4; 18pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             72.0%;
85.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
Matches 6; Conserv
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Gaps
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Score 36; DB 23; Length 8;
Pred. No. 9.3e+05;
0; Mismatches 1; Indels
                                6; Conservative
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Gaps

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Indels

6; Conservative

WSYWLRP 8 1 WSYGLRP 7

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RESULT 14 AAU76988

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treating breast cancer, dedometriosis, grometrium tumour, Alzheimer's disease, circulatory system disorders, menopausal disorders, irregular periods, cancer metastasis, premenstrual syndrome, muscular distress or osteopathies due to calcium/phosphorus imbalance. (I) comprises a gonadotrophin releasing hormone (GRRH) agonist and a selective oestrogen receptor modulator (SERM). (I) has cytostatic, gynaccological, nootropic, neuroprotective, cardiovascular, endocrine, muscular and osteopathic activities. (I) can be used as GMRH agonists used in combination with selective oestrogen receptor modulators, selective androgen receptor modulators, sex hormone synthesis inhibitors, lyase inhibitors, receptor-type tyrosine kinase inhibitors, bone metabolism requiators, antagonists for treating and preventing endometriosis, myometrium tumour, antagonists for treating and preventing endometriosis, myometrium tumour, irregular periods, cancer metastasis, premenstrual syndrome, muscular distress, osteopathies due to calcium/phosphorus imbalance,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    prostatomegally or prostate cancer or breast cancer or their reoccurrence or metastasis. Combination enhances the quality of life by enhancing GnRH agonist activity and/or reducing side effects. The present sequence represents a GnRH agonist related peptide which is given in the present
                                                                                                                                                                                                                                                                                                                       Gonadotrophin releasing hormone agonist; GnRH agonist; combination drug; pharmaceutical; breast cancer; endometriosis; myometrium tumour; Alzheimer's disease; circulatory system disorder; menopausal disorder; irregular period; cancer metastasis; premenstrual syndrome; osteopathy; muscular distress; calcium/phosphorus imbalance; SERM; gynaecological; selective oestrogen receptor modulator; cytostatic; nootropic; muscular; neuroprotective; cardiovascular; endocrine; osteopathic; prostatomegally;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Pharmaceutical composition e.g. for breast cancer comprises gonadotrophin releasing hormone agonist and selective estrogen receptor modulator
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention describes a pharmaceutical composition (I) for
                                                                                                                                                                                                                                                                                 Gonadotrophin releasing hormone agonist deslorelin peptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure; Page 8; 73pp; Japanese.
                                                                                                                                                   ABP96549 standard; peptide; 8 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     08-AUG-2002; 2002WO-JP08130.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (TAKE ) TAKEDA CHEM IND LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               10-AUG-2001; 2001JP-0244616
                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Furuya S, Kusaka M;
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WSYGLRP 7
                                      WSYWLRP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   8 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO2003015820-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 prostate cancer
                                                                                                                                                                                                                                        27-MAY-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            27-FEB-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           nvention.
                                                                                                                                                                                            ABP96549;
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DB 24; Length 8;

72.0%; Score 36; DB 24; 85.7%; Pred. No. 9.3e+05;

Query Match Best Local Similarity

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Gaps
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0
Indels
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Mismatches
                                                              completed: November 17, 2003, 18:40:33
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 6; Conservative
                  1 WSYGLRP
                                   2 WSYWLRP
                                                                       : 41 secs
 Matches
                                                                        Job time
                                                               Search
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5.1.6
Compugen Ltd.
GenCore version (c) 1993 - 2003
          Copyright
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OM protein - protein search, using sw model

November 17, 2003, 18:38:46; Search time 20 Seconds (without alignments) 38.467 Million cell updates/sec Run on:

US-09-462-089-3 50 Title: Perfect score:

WSYGLRPG 8 Scoring table: Sequence:

BLOSUM62 Gapop 10.0 , Gapext 0.5

283308 seqs, 96168682 residues Searched:

Total number of hits satisfying chosen parameters:

604

Minimum DB seq length: 0 Maximum DB seq length: 8

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

PIR 76: * 1: pir1: * 2: pir2: * 3: pir3: * 4: pir4: * Database :

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

STIMMARTES

	Description	fatty-acid synthas	neuropeptide calla	Ig H chain V-D-J r	Q.		244K exoantigen -	peptidyl-dipeptida	hypothetical TCL3	T-cell receptor be	acylase - Kluyvera	hypothetical prote	glucose-6-phosphat	leucokinin VIII -	Ig heavy chain CRD	RPCH-related neuro	Ig heavy chain CRD	ubiquitin - rat	photosystem I 10.4	Leu-enkephalin - b	Met-enkephalin - b	hypothetical prote	T-cell receptor be	T-cell receptor be	RNA-directed DNA p	halo-toxin - Pseud	T-cell receptor be	hypothetical colla	Met-enkephalin-Arg	platelet aggregati
SUMMARIES	QI	A60139	D47393	PH1618	155382	PT0281	A33098	JN0862	179564	PT0581	S19288	S16324	S11078	JS0318	PT0311	A34626	PT0240	847552	PQ0689	B61445	A61445	853595	PT0572	PT0714	A35890	A61049	PT0715	A35039	6022	A44428
	DB	2	~	7	4	~	~	7	4	~	7	~	~	~	~	~	Ģ	~	~	~	~	ď	~	7	~	7	7	4	~	7
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	Query Match	38.0	38.0	34.0				28.0	28.0	28.0			28.0			26.0	26.0	26.0	26.0	26.0								26.0		
	Score	19	19	17	16	15	15	14	14	14	14	14	14	14	14	13	13	13	13	13	13	13	13	13	13	13	13	13	13	13
	Result No.		7	m	4	5	9	7	80	σ	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	56	27	28	29

seed protein ws-5	dihydrofolate redu	Ig mu chain D regi	T-cell receptor be	T-cell receptor be	glutathione S-tran	red pigment-concen	adipokinetic hormo	adipokinetic hormo	adipokinetic hormo	unidentified 6.5/3	spasmogenic toxin	neuropeptide calla	T-cell receptor be	phenol 2-monooxyge	T-cell receptor be
E61491	148105	E33932	PT0515	PT0654	S66442	A61348	A33995	855310	A58620	PQ0701	A46306	E47393	PT0559	A37832	PT0661
7	~	~	7	7	7	~	~	7	7	7	7	7	7	~	N
7	7	7	7	7	7	8	œ	æ	œ	æ	œ	œ	œ	4	4
	0	0.	26.0	0.	26.0	26.0	26.0	26.0	26.0	26.0	26.0	26.0	6.0	4.0	24.0
26.0	26.0	56	56	56	56	3	Ñ	Ň	~	7	~	ñ	ā	~	N
13 26.0	13 26.	13 26	13 26	13 26	13 26	13 26	13 2	13 2	13 2	13 2	13 20	13 26	13 2	12 2	12 2

ALIGNMENTS

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fatty-acid synthase (EC 2.3.1.85) - rabbit (fragment)
Cispecies: Oryctolagus cuniculus (domestic rabbit)
Cjate: 22-Jan-1993 #sequence_revision 22-Jan-1993 #text_change 26-May-2000
CjAccession: A60139
RjHardie, D.G.; Dewart, K.B.; Altken, A.; McCarthy, A.D.
Biochim. Biophys. Acta 828, 380-382, 1985
A;Title: Amino acid sequence around the reactive serine residue of the thioesterase domc A;Reference number: A60139; MUID:85175165; PMID:3921056
A;Rocession: A60139
A;Rocession: A60139
A;Residues: 1-7 <HAR>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     C;Superfamily: rat fatty-acid synthase; 3-oxoacyl-[acyl-carrier-protein] synthase I homc ydrollase homology; Rort-chain alcohol dehydrogenase homology; [acyl-carrier-protein] S-C;Keywords: acyltransferase; carrier protein; coenzyme A; homodimer; multifunctional env F;S/Active site: Ser (of oleoyl-carrier-protein) hydrolase) #status experimental
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ö
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Matches 3; Conservative
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RESULT 1
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:||| 4 YSYG 7 1 WSYG 4 g

RESULT 2 D47393

neuropeptide callatostatin 4 - bluebottle fly (Calliphora vomitoria)

C;Species: Calliphora vomitoria C;Date: 16-Feb-1994 #sequence_revision 18-Nov-1994 #text_change 28-Apr-1995 C;Accession: D47393 R;Duve, H.; Johnsen, A.H.; Scott, A.G.; Yu, C.G.; Yagi, K.J.; Tobe, S.S.; Thorpe, A. Proc. Natl. Acad. Sci. U.S.A. 90, 2456-2460, 1993 A;Title: Callatostatins: neuropeptides from the blowfly Calliphora vomitoria with sequer A;Reference number: A47393; MUID:93211980; PMID:8460157

A,Accession: D47393
A,Status: preliminary
A,Molecule type: protein
A,Residues: 1-8 < DUV>
A,Experimental source: thoracic ganglia
A,Experimentel source thoracic ganglia

ö 38.0%; Score 19; DB 2; Length 8; 60.0%; Pred. No. 2.8e+05; tive 2; Mismatches 0; Indels Query Match
Best Local Similarity 60.0
Matches 3; Conservative

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Gaps

1 WSYGL 5 : |: | | 4 YSFGL g ò

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peptidyl-dipeptidase A inhibitory peptide C112 - striped bonito
C;Species: Sarda orientalis (striped bonito)
C;Date: 10-Mar-1994 #sequence_revision 10-Mar-1994 #text_change 07-May-1999
C;Accession: JN0862
B;Matsumura, N.; Fujii, M.; Takeda, Y.; Shimizu, T.
Biosci. Biotechnol. Biochem. 57, 1743-1744, 1993
A;Title: Isolation and characterization of angiotensin I-converting enzyme inhibitory px
A;Reference number: JN0865; MUID:94080036; PMID:7764272
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A;Molecule type: protein
A;Residues: 1-5 <MAT>
A;Residues: 1-5 <MAT>
A;Residues: 1-5 <MAT>
C;Comment: The amino terminal tripeptide of this protein inhibits angiotensin I-convert: C;Superfamily: bradykinin-potentiating peptide
C;Keywords: angiotensin-converting enzyme inhibitor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CyAccession: 179564

Rizutter, M.; Hockett, R.D.; Roberts, C.W.; McGuire, B.A.; Bloomstone, J.; Morton, C.C..

Brizutter, M.; Hockett, R.D.; Roberts, C.W.; McGuire, B.A.; Bloomstone, J.; Morton, C.C..

Proc. Natl. Acad. Sci. U.S.A. 87, 3161-3165, 1990

A;Title: The t(10;14) (q24;q11) of T-cell acute lymphoblastic leukemia juxtaposes the della; Reference number: 159162; MUID: 90222189; PMID: 2326274

A;Reference number: 159162; MUID: 90222189; PMID: 2326274

A;Accession: 179564

A;Accession: 179564

A;Accession: Los Aire

A;Residues: 1-6 <2UT>
A;Residues: 1-6 <2UT>
A;Cross-references: GB:M33602; NID: 9339907; PIDN: AAA66449.1; PID: 9807656
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                                                                                                                                                                                                                                                                                                                                                         C;Species: Plasmodium falciparum
C;Date: 24-Aug-1990 #text_change 09-Jun-2000
C;Accession: A33098
R;Nichols, J.H; Hager, L.P.
A;Reference number: A33098
A;Reference number: A33098
A;Accession: A33098
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          C;Species: Homo sapiens (man)
C;Date: 28-Jan-2000 #sequence_revision 28-Jan-2000 #text_change 28-Jan-2000
                                   Gaps
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                                                                                                                                                                                                                                                                                                                                              244K exoantigen - malaria parasite (Plasmodium falciparum) (fragments)
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Pred. No. 2.8e+05;
1; Mismatches 0; Indels
                                   Indels
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       8e+05;
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Pred. No. 2.8e+05;
Similarity 100.0%; Pred. No. 2.8
2; Conservative 0; Mismatches
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Best Local Similarity 75.0%;
Matches 3; Conservative
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Best Local Similarity 66.7%;
Matches 2; Conservative
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A,Molecule type: protein
A,Residues: 1-7 <NIC>
Best Local Similarity
Matches 2; Conserv
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1 IRP 3
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R;Dawson, S.J.; Wiman, B.; Hamsten, A.; Green, F.; Humphries, S.; Henney, A.M.
J. Baiol. Chem. 288, 10739-10745, 1993
A;Itle: The two allele sequences of a common polymorphism in the promoter of the plasmin A;Reference number: 155382
A;Accession: 155382
A;Accession: 155382
A;Accession: 155382
A;Cross-references: GB:M91557; NID:g190020; PIDN:AAA60110.1; PID:g190021
A;Residues: 1-7 cDAW
A;Residues: 3 common control 
                                                                                                                                                                                                 C,Accession: PH1618
R;Levinson, D.A.; Campos-Torres, J.; Leder, P.
J. Exp. Med. 178, 317-329, 1993
A;Title: Molecular characterization of transgene-induced immunodeficiency in B-less mice A;Reference number: PH1580; MUID:93301609; PMID:8315387
A;Accession: PH1618
A;Molecula type: DNA
A;Residues: 1-8 <LEV>A;Residues: 1-8 <LEV>A;Residues: 1-8 <LEV>A;Residues: 1-8 <LEV>A;Residues: immunoglobulin
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[; Species: Homo sapiens (man)
[; Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change 16-Aug-1996
[; Accession: Pr0281
R; Yamada, M.; Wasserman, R.; Reichard, B.A.; Shane, S.; Caton, A.J.; Rovera, G. A.J. Exp. Med. 173, 395-407, 1991
A; Title: Preferential utilization of specific immunoglobulin heavy chain diversity and A; Reference number: Pr0222; MUID:91108337; PMID:1899102
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C;Date: 16-Apr-1999 #sequence_revision 16-Apr-1999 #text_change 20-Apr-2000
C;Accession: 155382
                                                                                                      1 H chain V-D-J region (clone B-less 33) - mouse (fragment)
Species: Mus musculus (house mouse)
Date: 02-Jun-1994 #sequence_revision 02-Jun-1994 #text_change 17-Mar-1999
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Pred. No. 2.8e+05;
1; Mismatches 1;
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A;Residues: 1-5 < YAM>
A;Experimental source: B lymphocyte
C;Keywopds: heterotetramer; immunoglobulin
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50.0%;
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Best Local Similarity 50.0°
---- 2; Conservative
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Matches 3; Conservative
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C;Accession: S16324
R;Ruberti, I.; Sessa, G.; Lucchetti, S.; Morelli, G.
EMBO J. 10, 1787-1791, 1991
A;Title: A novel class of plant proteins containing a homeodomain with a closely linked A;Reference number: S16323; MUID:91266907; PMID:1675603
A;Accession: S16324
A;Status: translation not shown
C; Comment: This is the hypothetical translation of a sequence translated in an incorrect
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        C;Species: Kluyvera cryocrescens
C;Species: Kluyvera cryocrescens
C;Date: 19-Mar-1997 #sequence_revision 19-Mar-1997 #text_change 24-Jul-1997
C;Date: 19-Mar-1997 #sequence_revision 19-Mar-1997 #text_change 24-Jul-1997
C;Caccession: S19288
R;Martin, J; Slade, A; Aitken, A; Arche, R; Virden, R.
Biochem, J. 280, 659-662, 1991
Biochem, J. 280, 659-662, 1991
A;Title: Chemical modification of serine at the active site of penicillin acylase from 1A;Reference number: S19288 MUID:92109664; PMID:1764029
A;Accession: S19288
A;Accession: S19288
A;Accession: S19288
A;Accession: S19288
A;Accession: S19288
A;Residues: 1-8 cMAR>
                                                                                                                                                                                                                                                                                                                                                                                                                         J. Exp. Med. 174, 115-124, 1991
A/Title: Junctional sequences of fetal T cell receptor beta chains have few N regions.
A/Reference number: PT0509; MUID:91277601; PMID:1711558
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C;Species: Arabidopsis thaliana (mouse-ear cress)
C;Date: 21-Nov-1993 #sequence_revision 12-May-1995 #text_change 21-Jul-2000
                                                                                                                                                                                                                                                                                                                  -cell receptor beta chain V-D-J region (159-1A) - mouse (fragment)
:Species: Mus musculus (house mouse)
;Date: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 30-May-1997
;Accession: PT0581
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Pred. No. 2.8e+05;
                                               Score 14; DB 4; Length 6;
Pred. No. 2.8e+05;
0; Mismatches 2; Indels
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A;Molecule type: mRNA
A;Residues: 1-7 <FEE>
A;Experimental source: day 19 fetal thymus, strain BALB/c
C;Keywords: T-cell receptor
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Best Local Similarity 75.0%;
Matches 3; Conservative (
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50.0%;
                                               Query Match
Best Local Similarity 50.0%;
Matches 2; Conservative
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Best Local Similarity 50.0
Matches 2; Conservative
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R, Feeney, A.J.
J. Exp. Med. 17
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A;Title: Fast atom bombardment mass spectrometry and chemical analysis in determination. A;Reference number: $11074; MUID:90353571; PMID:2387402
A;Accession: $11078
A;Accession: $11078
A;Molecule type: protein
A;Residues: 1-8 <EGE>
A;Note: the source is designated as Pichia jadinii
C;Keywords: acetylated amino end; oxidoreductase; pentose phosphate pathway
F;1/Modified site: acetylated amino end (Ser) #status experimental
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C;Species: Leucophaea maderae (Madeira cockroach)
C;Species: Leucophaea maderae (Madeira cockroach)
C;Date: 07-Sep-1990 #sequence_revision 07-Sep-1990 #text_change 20-Jun-2000
C;Date: 07-Sep-1990 #sequence_revision 07-Sep-1990 #text_change 20-Jun-2000
C;Accession: JSO318
R;Holman, G.M.; Cook, B.J.; Nachman, R.J.
Comp. Biochem. Physiol. C 88, 31-34, 1987
A;Title: Isolation, primary structure and synthesis of leucokinins VII and VIII: the fix A;Reference number: JSO317
A;Accession: JSO318
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      R;Egestad, B.; Estonius, M.; Danielsson, O.; Persson, B.; Cederlund, E.; Kaiser, R.; Hol
FEBS Lett. 269, 194–196, 1990
                                                                                                                                                                                                                                                                                                                                                                                                                      glucose-6-phosphate 1-dehydrogenase (EC 1.1.1.49) - yeast (Pichia jadinii) (fragment)
C;Species: Pichia jadinii, Candida utilis
C;Date: 30-Jun-1991 #sequence_revision 30-Sep-1991 #text_change 05-Aug-1994
C;Accession: S11078
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C;Species: Homo sapiens (man)
C;Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change 16-Aug-1996
C;Accession: PT0311
R;Yamada, M.; Wasserman, R.; Reichard, B.A.; Shane, S.; Caton, A.J.; Rovera, G.J. Exp. Med. 173, 395-407, 1991
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A,Molecule type: mRNA
A,Residues: 1-8 <RUB>
A,Cross-references: EMBL:X58821; NID:g16327; PIDN:CAA41624.1; PID:g579259
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Pred. No. 2.8e+05;
2; Mismatches 0; Indels
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Similarity 50.0%;
2; Conservative
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Best Local Similarity 60.0°
Matches 3; Conservative
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PT0311
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RPCH-related neuropeptide - ferruginous spindle
C;Species: Fusinus ferrugineus (ferruginous spindle)
C;Species: Fusinus ferrugineus (ferruginous spindle)
C;Date: 06-Jul-1990 #sequence_revision 06-Jul-1990 #text_change 31-Dec-1993
C;Accession: A34666
R;Kuroki, Y.; Kanda, T.; Kubota, I.; Fujisawa, Y.; Ikeda, T.; Miura, A.; Minamitake, Y.;
Biochem. Biophys. Res. Commun. 167, 273-279, 1990
A;Title: A molluscan neuropeptide related to the crustacean hormone, RPCH.
A;Reference number: A34626; MUID:90179762; PMID:2310394
A,Title: Preferential utilization of specific immunoglobulin heavy chain diversity and A;Reference number: PT0222; MUID:91108337; PMID:1899102
A;Accession: PT0311
A;Molecule type: DNA
A;Residues: 1-8 <YAM>
A;Experimental source: B lymphocyte
C;Keywords: heterotetramer; immunoglobulin
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28.0%; Score 14; DB 2; Length 8;
Best Local Similarity 66.7%; Pred. No. 2.8e+05;
Matches 2; Conservative 1; Mismatches 0; Indels
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A,Molecule type: protein
A,Residues: 1-4 <KUR>
C,Keywords: neuropeptide
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3 AYG 5
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Search completed: November 17, 2003, 18:42:15 Job time : 20 secs

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[1]
SEQUENCE.
TISSUE=Cerebral ganglion, and Thoracic ganglion;
MEDLINE=98121193; Pubmed=9461295;
Duve H., Johnsen A.H., Maestro J.-L., Scott A.G., Jaros P.P.,
Thorpe A.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RAKKR
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P81817 carcinus ma
P81818 carcinus ma
P81821 carcinus ma
P81182 carcinus ma
P82154 cydia pomon
P81185 cydia pomon
P81180 carcinus ma
P8180 carcinus ma
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P81815 cydia pomon
P81152 cydia pomon
P82157 cydia pomon
P81181 carcinus ma
P8181 carcinus ma
P8184 pinus pinas
P8184 mus musculu
P3009 homo sapien
P8180 herpes simp
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                                                                              November 17, 2003, 18:32:45; Search time 11 Seconds (without alignments) 34.201 Million cell updates/sec
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P21140
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GenCore version 5.1.6
Copyright (c) 1993 - 2003 Compugen Ltd.
                                                                                                                                                                                                                                                             Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                      127863 seqs, 47026705 residues
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Maximum Match 100%
Listing first 45 summaries
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Gapop 10.0 , Gapext 0.5
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Perfect score:
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LCK4_LEUMA LCK5_LEUMA LCK6_LEUMA LCK6_LEUMA LCK7_LEUMA	PRI; 8 AA. eated) ist sequence update) ist annotation update) shore crab) (Green crab). Troppoda; Crustacea; Malacostraca; i; Decapoda; Crustacea; Malacostraca; i; Portunidae; Carcinus. i, and Thoracic ganglion; le9461295; lestro JL., Scott A.G., Jaros P.P., action of multiple neuropeptides of the in the shore crab Carcinus maenas."; -734(1997). TO THE ALLATOSTATIN FAMILY. Multigene family. Multigene family. i) 7C286845AA476878 CRC64; i) 7C286845AA4476878 CRC64; i) 7C286845AA4476878 CRC64; i) 7L286ASAA446978 CRC64; ii) Mismatches 0; Indels 0; Gaps 0;	PRT; 5 AA. pence update) totation update) crab) (Green crab). trustaces, Malacostraca; proda; Pleocyemata; Brachyura; unidae; Carcinus.
34 12 24.0 8 1 LCK5 35 12 24.0 8 1 LCK5 36 12 24.0 8 1 LCK7 37 12 24.0 8 1 LCK7 39 11 22.0 4 1 OCP3 40 11 22.0 6 1 EF01 42 11 22.0 7 1 BRHP 44 11 22.0 7 1 TPFY 45 11 22.0 7 1 TPFY	RESULT 1 ALI6_CARMA 1D ALI6_CARMA ALS GLARMA STANDARD; PRT; R M AU 30-MAY-2000 (Rel. 39, Created) DT 30-MAY-2000 (Rel. 39, Last sequence update) Carcinus maenas (Common shore crab) (Green Eumalacostraca; Eucarida; Decapoda; Pleocyc NEMILARIDARIO PROMENT PROTUNIDAS, Portunidae; Carci RX MEDLINE-88121193; Pubméd-946125; RA Thorpe A; Duve H., Johnsen A.H., Maestro JL., Scott RA Thorpe A; TISOLation and identification of multiple rangement of the shore crab RY Allatostatin superfamily in the Shore crab RY Neuropeptide; Amidation; Multigene family. RY Neuropeptide; Score 23; DB Best Local Similarity 80.0%; Pred. No. 1.38 Matches 4; Conservative 1; Mismatches	QY 1 WSYGL 5 Db

us-09-462-089-3.closed.rsp

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TISSUE=Larva;
MEDLINE=98054539; PubMed=9392829;
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NCBI_TaxID=82600;
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-!- FUNCTION: MAY ACT AS A NEUROTRANSMITTER OR NEUROMODULATOR.
-!- SIMILARITY: BELONGS TO THE ALLATOSTATIN FAMILY.
Neuropeptide; Amidation; Multipene family.
"Isolation and identification of multiple neuropeptides of the
            allatostatin superfamily in the shore crab Carcinus maenas.";
Eur. J. Biochem. 250:727-734(1997).
-!- FUNCTION: MAY ACT AS A NEUROTRANSMITTER OR NEUROMODULATOR.
-!- SIMILARITY: BELONGS TO THE ALLATOSTATIN FAMILY.
MOD RES

S AMIDATION (POTENTIAL).
SEQÜENCE 5 AA; 586 MW; 672879D5AB300000 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Duve H., Johnsen A.H., Maestro J.-L., Scott A.G., Jaros P.P.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Carcinus maenas (Common shore crab) (Green crab).
Bukaryota; Metazoa, Arthropoda; Crustacea; Malacostraca;
Eumalacostraca; Bucarida; Decapoda; Pleccyemata; Brachyura;
Bubrachyura; Portunoidea; Portunidae; Carcinus.
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                                                                                                                                                                                Score 19; DB 1; Length 5;
Pred. No. 1.3e+05;
2; Mismatches 0; Indels
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MEDLINE=98121193; PubMed=9461295;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TISSUE=Cerebral ganglion, and Thoracic ganglion; MEDLINE=98121193; PubMed=9461295;
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30-MAY-2000 (Rel. 39, Last sequence update)
30-MAY-2000 (Rel. 39, Last annotation update)
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30-MAY-2000 (Rel. 39, Created)
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60.0%;
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Gaps
"Isolation and identification of multiple neuropeptides of the allatostatin superfamily in the shore crab Carcinus maenas."; Eur. J. Biochem. 250:727-734(1997).
-!- FUNCTION: MAY ACT AS A NEUROTRANSMITTER OR NEUROMODULATOR.
-!- SIMILARITY: BELONGS TO THE ALLATOSTATIN FAMILY.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 allatostatin superfamily in the shore crab Carcinus maenas.";
Eur. J. Blochem. 250:727-734(1997).
-!- FUNCTION: MAY ACT AS A NEUROTRANSMITTER OR NEUROMODULATOR.
-!- SIMILARITY: BELONGS TO THE ALLATOSTATIN FAMILY.
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Bukaryota; Metazoa, Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Lepidoptera; Glossata; Ditrysia;
Tortricoidea; Tortricidae; Olethreutinae; Cydia.
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Eumalacostraca; Eucarida; Decapoda; Pleocyemata; Brachyura;
Eubrachyura; Portunoidea; Portunidae; Carcinus.
                                                                                                                                                                                                                                                                                  Length 8;
                                                                                                                                                                                                                                                                                                                                            Indels
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MOD RES 8 AMIDATION (POTENTIAL)
SEQUENCE 8 AA, 919 MW; C82879D5AB569AB5 CRC64;
                                                                                                                                                         Neuropeptide, Amidation, Multigene family.

MOD RES 8 8 AMIDATION (POTENTIAL)

SEQUENCE 8 AA, 858 MW; C82879D5AB46D865 CRC64;
                                                                                                                                                                                                                                               38.0%; Score 19; DB 1; Ler
60.0%; Pred. No. 1.38+05;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TIŜSUE-Cerebral ganglion, and Thoracic ganglion;
MEDLINE-98121193; PubMed-9461295;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (Rel. 39, Created)
(Rel. 39, Last sequence update)
(Rel. 39, Last annotation update)
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Last annotation update)
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30-MAY-2000 (Rel. 39, Last seq
30-MAY-2000 (Rel. 39, Last anno
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8 AA.

PRT;

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STANDARD;
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ALL4 CYDPO
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ALL2 CARMA
LID ALLS CARMA
LID 30-MAY
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Matches
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"Distribution and functional significance of Leu-callatostatins in "Distribution and functional";

[the blowfly Calliphora vomitoria",

[cell Tissue Res. 276:367-379(1994)]

-!- FUNCTION: MAY ACT AS A NEUROTRANSHITTER OR NEUROMODULATOR AND PLAY A ROLE IN THE INTEGRATION OF INFORMATION WITHIN THE BRAIN. MAY BE INVOLVED IN THE COMTROL OF VISCERAL MUSCLES DUE TO ITS ABILITY TO BEHAVE AS POTENT INHIBITORS OF PERISTALIC MOVEMENTS. MAY ALSO FULFILL A NEUROPROMONAL ROLE ON WISCLES OF THE GUT AND HEART.

-!- TISSUE SPECIFICITY: BRAIN, SUBORESOPHAGEAL GANGLION, RETROCEREBRAL COMPLEX. THORACICO-ABDOMINAL GANGLION, PERIPHERAL NEUROSECRETORY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Calliphora vomitoria (Blue blowfly).
Bukaryota, Metazoa, Arthropoda, Hexapoda, Insecta, Pterygota;
Neopiera, Endopterygota; Diptera, Brachycera, Muscomorpha, Oestroidea;
Calliphoridae; Calliphora.
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Proc. Natl. Acad. Sci. U.S.A. 90:2456-2460(1993).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDLINE=93211980; PubMed=8460157;
Duve H., Johnsen A.H., Scott A.G., Yu C.G., Yagi K.J., Tobe S.S.,
Duve H., Johnsen A.H., Maestro J.-L., Scott A.G., Winstanley D.
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                   Davey M., East P.D., Thorpe A.;
"Lepidopteran peptides of the allatostatin superfamily.";
"Peptides 18:130-1309 (1997)
-!- SIMILARITY: BELONGS TO THE ALLATOSTATIN FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              38.0%; Score 19; DB 1; Length 8; 60.0%; Pred. No. 1.3e+05; tive 2; Mismatches 0; Indels
                                                                                                                                                                                                    Similarity 60.0%; Pred. No. 1.3e+05;
3; Conservative 2; Mismatches 0; Indels
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                                                                                                                                                            8 AA; 926 MW; C82879D5AB477415 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-NOV-1995 (Rel. 32, Created)
01-NOV-1995 (Rel. 32, Last sequence update)
30-MAY-2000 (Rel. 39, Last annotation update)
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                                                                                                                                      AMIDATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CHARACTERIZATION.
MEDLINE=94291167; PubMed=8020069;
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Best Local Similarity 60.0
Matches 3; Conservative
                                                                                                         Neuropeptide; Amidation.
MOD RES 8 8
SEQUENCE 8 AA; 926 MW;
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MOD_RES 8 8
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Best Local Similarity
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ID ALL4 CALVO
AC P41840;
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RESULT 8 ALL4_CYDPO

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-!- FUNCTION: MAY ACT AS A NEUROTRANISMITTER OR NEUROMODULATOR.
-!- SIMILARITY: BELONGS TO THE ALLATOSTATIN FAMILY.
                                                                        Cydiastatin 4.
Cydia pomonella (Codling moth).
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Lepidoptera; Glossata; Ditrysia;
Tortricoidea; Tortricidae; Olethreutinae; Cydia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Duve H., Johnsen A.H., Maestro J.-L., Scott A.G., Jaros P.P.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Carcinus means (Common shore crab) (Green crab).

Eukaryota; Metazoa; Arthropoda; Crustacea; Malacostraca;

Eumalacostraca; Eucarida; Decapoda; Pleocyemata; Brachyura;

Eubrachyura; Portunoidea; Portunidae; Carcinus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  38.0%; Score 19; DB 1; Length 8; 60.0%; Pred. No. 1.3e+05; tive 2; Mismatches 0; Indels
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P81805;
30-MAY-2000 (Rel. 39, Created)
30-MAY-2000 (Rel. 39, Last sequence update)
30-MAY-2000 (Rel. 39, Last annotation update)
30-MAY-2000 (Rel. 39, Created)
30-MAY-2000 (Rel. 39, Last sequence update)
30-MAY-2000 (Rel. 39, Last annotation update)
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MOD RES 7 7 AMIDATION (PO
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4 YSFGL 8
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ALL3 CARMA
ID ALL3_CARMA
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                                                                                                                                                                                                                                                  "Isolation and identification of multiple neuropeptides of the allatostatin superfamily in the shore crab Carcinus maenas."; Eur. J. Biochem. 250:727-734 (1997).

-i. FUNCTION: MAY ACT AS A NEUROTRANSMITTER OR NEUROMODULATOR.-i. SIMILARITY: BELONGS TO THE ALLATOSTATIN FAMILY.

MOLYOPEPLIDE: Amidation; Multigene family.

AMIDATION.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             allatostatin superfamily in the shore crab Carcinus maenas.";
Eur. J. Biochem. 250:727-734(1997).
-!- FUNCTION: MAY ACT AS A NEGROTRANSMITTER OR NEUROMODULATOR.
-!- SIMILARITY: BELONGS TO THE ALLATOSTATIN FAMILY.
Neuropeptide; Multigene family.
SEQUENCE 8 AA; 913 MW; 672879CDCB569AB7 CRC64;
                                                                                                                                                                                                                     Duve H., Johnsen A.H., Maestro J.-L., Scott A.G., Jaros P.P.,
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Carcinus maenas (Common shore crab) (Green crab).

Bukaryota, Metazoa; Arthropoda; Crustacea; Malacostraca;

Bunalacostraca; Bucarida; Decapoda; Pleocyemata; Brachyura;

Bubrachyura; Portunoidea; Portunidae; Carcinus.
                                                             Carcinus maenas (Common shore crab) (Green crab).
Eukaryota; Metazoa; Arthropoda; Crustacea; Malacostraca;
Eumalacostraca; Eucarida; Decapoda; Pleocyemata; Brachyura;
Eubrachyura; Portunoidea; Portunidae; Carcinus.
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                                                                                                                                                                                                                                                                                                                                                                       7 AA; 781 MW; 672879CDCB476420 CRC64;
                                                                                                                                                                                TISSUE=Cerebral ganglion, and Thoracic ganglion; MEDLINE=98121193; PubMed=9461295;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          30-MAY-2000 (Rel. 39, Created)
30-MAY-2000 (Rel. 39, Last sequence update)
30-MAY-2000 (Rel. 39, Last annotation update)
30-MAY-2000 (Rel. 39, Created)
30-MAY-2000 (Rel. 39, Last sequence update)
30-MAY-2000 (Rel. 39, Last annotation update)
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                                                                                                                                                                                                                                                                                                                                                                                                                                          3; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                          2; Conservative
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3 YAFGL 7
                                                                                                                                    NCBI_TaxID=6759;
                                                    Carcinustatin 5
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P82152;
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                                                                                                                                                                                                                                      Thorpe A.;
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                                                                                                                                                                                                                                                                    "Isolation and identification of multiple neuropeptides of the allacotatin superfamily in the shore crab Carcinus maenas."; Eur. J. Biochem. 250:727-734 (1997).
-!- FUNCTION: MAY ACT AS A NEUROTRANSMITTER OR NEUROMODULATOR.
-!- STMILARMITY: BELONGS TO THE ALLATOSTATIN FAMILY.
Neuropeptide; Multigene family.
SEQUENCE 7 AA; 796 MW; 672879CDCB476B70 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                          ;
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-!- FUNCTION: MAY ARACT AS A NEUROTRANSMITTER OR NEUROMODULATOR.
-!- SIMILARITY: BELONGS TO THE ALLATOSTATIN FAMILY.
Neuropeptide; Multigene family.
                                                                                                                                                                                     TISSUE=Cerebral ganglion, and Thoracic ganglion;
MEDLINE=98121193; PubMed=9461295;
Duve H., Johnsen A.H., Maestro J.-L., Scott A.G., Jaros P.P.,
Thorpe A.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Duve H., Johnsen A.H., Maestro J.-L., Scott A.G., Jaros P.P.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Carcinus maenas (Common shore crab) (Green crab).
Eukaryota; Metazoa; Arthropoda; Crustacea; Malacostraca;
Eumalacostraca; Eucarida; Decapoda; Pleocyemata; Brachyura;
Eubrachyura; Portunoidea; Portunidae; Carcinus.
                                                                                  Carcinus maenas (Common shore crab) (Green crab).
Bukaryota; Metazoa; Arthropoda; Crustacea; Malacostraca;
Eumalacostraca; Bocarida; Decapoda; Pleccyemata; Brachyura;
Bubrachyura; Portunoidea; Portunidae; Carcinus.
                                                                                                                                                                                                                                                                                                                                                                                                       Score 16; Lu -.
Pred. No. 1.3e+05;
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                                                                                                                                                                                                                                                                                                                                                                                                         32.0%; Score 16; DB 1; Length 7; 40.0%; Pred. No. 1.3e+05;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TISSUE-Cerebral ganglion, and Thoracic ganglion; MEDLINE-98121193; PubMed-9461295;
             30-MAY-2000 (Rel. 39, Created)
30-MAY-2000 (Rel. 39, Last sequence update)
30-MAY-2000 (Rel. 39, Last annotation update)
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30-MAY-2000 (Rel. 39, Last sequence update)
30-MAY-2000 (Rel. 39, Last annotation update)
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Matches 2; Conservative
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Matches 2; Conserv
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3 YAFGL 7
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                                                                    Carcinustatin 3.
                                                                                                                                                      NCBI_TaxID=6759;
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ID ALLS CARMA
AC P81808;
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ALL4_CARMA
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TISSUB-Learva;

MEDLINE-98054539; PubMed-9392829;

MEDLINE-98054539; PubMed-9392829;

Duve H., Johnsen A.H., Masstro J.-L., Scott A.G., Winstanley D.,

Davey M., East P.D., Thorpe A.;

"Lepidopteran peptides of the allatostatin superfamily.";

"Eptides 18:1301-1309(1997).

-!- SIMILARITY: BELONGS TO THE ALLATOSTATIN FAMILY.

Neuropeptide; Amidation.

MOD RES

AMIDATION.
                                                                                                                              .;
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Cydia pomonella (Codling moth).
Eukaryota metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Bukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Booptera; Endopterygota; Lepidoptera; Glossata; Ditrysia;
Tortricoidea; Tortricidae; Olethreutinae; Cydia.
NCBI_TaxID=82600;
                                                   Cydia pomonella (Codling moth).
Bukaryota, Metazoa, Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera, Endopterygota; Lepidoptera; Glossata; Ditrysia;
Tortricoidea; Tortricidae; Olethreutinae; Cydia.
                                                                                                                                                                                                                                                                                          32.0%; Score 16; DB 1; Length 8; 40.0%; Pred. No. 1.3e+05; Live 3; Mismatches 0; Indels
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8 AA; 934 MW; C82879C45B51F775 CRC64;
30-MAY-2000 (Rel. 39, Created)
30-MAY-2000 (Rel. 39, Last sequence update)
30-MAY-2000 (Rel. 39, Last annotation update)
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30-MAY-2000 (Rel. 39, Last sequence update)
30-MAY-2000 (Rel. 39, Last annotation update)
Cydiastatin 6.
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Best Local Similarity 40.0
Matches 2; Conservative
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Matches 2; Conservative
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4 YNFGL 8
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4 YNFGL 8
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                                         Cydiastatin 1
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SEQUENCE
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ALL6_CYDPO
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Search completed: November 17, 2003, 18:40:56 Job time : 11 secs

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Title: Perfect score:

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Scoring table: Sequence:

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Q47555 secherichia
089100 borrelia bu
088100 borrelia bu
088100 borrelia ol
808100 sepien
089100 sepien
82685 periplaneta
82687 periplaneta
82687 periplaneta
82689 periplaneta
82680 seriplaneta
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8260 
                                                  Q9tt78 canis famil
Q9xsyl canis famil
Q9szl3 oryctolagus
Q81802 zea mays (m
P82079 limnodynast
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lytechinus
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Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Lepidosauria; Squamata; Scleroglossa; Anguimorpha; Varanidae; Varanus.
NCBI_TaxID=169851;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ast J.C.; "Micochondrial DNA evidence and evolution in Varanoidea (Squamata)."; Cladistics 17:0-0(2001). BMBL; AF407521; AAL10116.1; -... Mitochondrion.
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2; Conservative 1; Mismatches 0; Indels
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01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
Cytochrome c oxidase subunit I (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               8 8 AA; 1053 MW; FE2729D5A36411A6 CRC64;
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P79940
PT PT; 8 AA.

AC P79940;
DT 01-MAY-1997 (TrEMBLrel. 03, Created)
DT 01-MAY-1997 (TrEMBLrel. 03, Last sequence update)
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09XSY1
095213
08L802
P82079
047505
098866
052062
Q15901
O02032
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Q81VK3
P82686
P82687
P82687
P826887
P82688
Q9GMH3
Q9GMH3
Q994VF
Q94VB
Q94VJ
Q94VJ
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                                      PRELIMINARY;
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   Query Match
Best Local S:
Matches 2;
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SEQUENCE
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Q94VC1
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O49223 glycine max
O09258 synechococc
O56140 streptococc
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                                                                                                                         November 17, 2003, 18:38:10 ; Search time 33 Seconds (without alignments) 62.558 Million cell updates/sec
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                   GenCore version 5.1.6
Copyright (c) 1993 - 2003 Compugen Ltd.
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Gapop 10.0 , Gapext 0.5
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sp_mammal:*
sp_mhc:*
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Maximum Match 100%
Listing first 45 sv
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Result Š.

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STRAIN'66, 71, 75, 90, 1, and 2;
Randi E., Pierpaoli M., Beaumont M., Ragni B., Sforzi A.;
Randi E., Pierpaoli M., Beaumont M., Ragni B., Sforzi A.;
"Genetic identification of wild and domestic cats (Felis silvestris),
and their hybrids using Bayesian clustering methods.";
Submitted (MAR-2001) to the EMBL/GenBank/DDBJ databases.
BMBL; AJ409137; CAC41051.1;
EMBL; AJ409138; CAC41057.1;
EMBL; AJ409141; CAC41060.1;
EMBL; AJ409141; CAC41060.1;
EMBL; AJ409141; CAC4106.1;
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Lepidosauria, Squamata, Scleroglossa, Anguimorpha, Varanidae, Varanus
NCBI_TaxID=169832;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ast J.C.;
"Mitochondrial DNA evidence and evolution in Varanoidea (Squamata).";
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                                    Eukaryota; Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Carnivora, Fissipedia, Felidae, Felis.
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01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
Cytochrome c oxidase subunit II (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            01-DEC-2001 (TrEMBLrel. 19, Created)
01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
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EMBL; AF407524; AAL10125.1; -.
Mitochondrion.
Felis silvestris (Wild cat).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Varanus salvator togianus. Mitochondrion.
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                                                                                                                  SEQUENCE FROM N.A.
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SEQUENCE FROM N.A.
                                                                              NCBI_TaxID=9683;
                                                                                                                                                                                                                                                                                                                                                      Mitochondrion.
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SEQUENCE
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Q94VB2;
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Q94PX7
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            XMeis1-4 protein (Fragment).
Xenopus laevis (African clawed frog).
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Amphibia; Batrachia; Anura; Mesobatrachia; Pipoidea; Pipidae;
Xenopodinae; Xenopus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens (Human).
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Caskey C.T.H.;
"Isolation of chromosome-specific genes by reciprocal probing of arrayed cDNAs and cosmid libraries.";
Hum. Mol. Genet. 0:0-0(1995).
EMBL; L32069; AAA73878.1; -.
                                                                                                                                                                            MEDLINE=97202105; PubMed=9049632; Steelman S., Moskow J.J., Muzynski K., North C., Druck T., Moskow J.C., Muebner K., Daar I.O., Buchberg A.M.; Montgomery J.C., Huebner K., Daar I.O., Buchberg A.M.; Identification of a conserved family of Mais1-related homeobox
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2; Conservative 0; Mismatches 1; Indels
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01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
Cytochrome c oxidase subunit II (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        01-NOV-1996 (TrEMBLrel. 01, Created)
01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
(Clone XPLSHBA) (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    8 AA; 1068 MW; 0315A37EAB5B0763 CRC64;
      01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
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AA.
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                                                                                                                                                                                                                                                                        Genome Res. 7:142-156(1997).
EMBL; U68389; AAB19199.1; -.
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Best Local Similarity
2; Conserva
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Best Local Similarity
Matches 2; Conserv
                                                                                                                                                          SEQUENCE FROM N.A.
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TISSUE=Placenta;
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Q15888;
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Vāranus salvator cumingi.
Mitochondrion.
Eukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Lepidosauria; Squamata; Scleroglossa; Anguimorpha; Varanidae; Varanus.
NCBI_TaxID=169830;
                                                                                                                                                                                                   Eukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Lepidosauria; Squamata; Scleroglossa; Anguimorpha; Varanidae; Varanus.
VCBI_TaxID=169831;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Agt J.C.;

"Mitochondrial DNA evidence and evolution in Varanoidea (Squamata).";

Cladistics 17:0-0(2001).

EMBL, AF407523; AAL10122.1; -.

Mitochondrion.

NON_TER
                                                                                                                                                                                                                                                                                                                                                       Ast J.C.;
"Mitochondrial DNA evidence and evolution in Varanoidea (Squamata).";
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01-OCT-2000 (TrEMBLrel. 15, Last sequence update)
01-MAR-2001 (TrEMBLrel. 16, Last annotation update)
38ba non-arginase growth inhibitory factor (NAGIF) (Fragment).
Rattus norvegicus (Ral).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
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Pred. No. 8.3e+05;
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                                                                                                       Cytochrome c oxidase subunit I (Fragment).
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                           Created)
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EMBL; AF407526; AAL10130.1; -.
Mitochondrion.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              8 AA; 992 MW;
                                                                                                                                                            Varanus salvator salvator
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Matches 2; Conservative
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Matches 2; Conservative
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                                                                                                                                                                                        Mitochondrion.
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P82598;
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P82598
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                                                                                                                                                                                  STRAIN=1, 2, 7, 12, 16, 17, and 110;
Randi E., Pierpaoli M., Beaumont M., Ragni B., Sforzi A.;
"Genetic identification of wild and domestic cats (Felis silvestris),
and their hybrids using Bayesian clustering methods.";
Submitted (MAR-2001) to the EMBL/GenBank/DDBJ databases.
EMBL, AJ409128; CAC41027.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE FROM N.A.

STRAIN=40, 1, 2, and 7;

Randi E., Pierpaoli M., Beaumont M., Ragni B., Sforzi A.;

Randi E., Pierpaoli M., Beaumont M., Ragni B., Sforzi A.;

"Genetic identification of wild and domestic cats (Felis silvestris),

and their hybrids using Bayesian clustering methods.";

Submitted (MAR-2001) to the EMBL/GenBank/DDBJ databases.

EMBL; AJ409140; CAC41048.1; -.

EMBL; AJ409144; CAC41069.1; -.

EMBL; AJ409144; CAC41069.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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                                                     Euteleostomi;
Felis.
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Mammalia, Eutheria, Carnivora, Fissipedia, Felidae, Felis.
NCBI_TaxID=61377,
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                                                  Eukaryota, Metazoa, Chordata, Craniata, Vertebrata,
Mammalia, Eutheria, Carnivora, Fissipedia, Felidae,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-DEC-2001 (TrEMBLrel. 19, Created)
01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
Cytochrome c oxidase subunit II (Fragment).
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8 AA; 951 MW; 262685BDC5A3733B CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                   AJ409131; CAC41036.1; -. AJ409132; CAC41039.1; -. AJ409133; CAC41042.1; -. AJ409134; CAC41045.1; -.
Felis silvestris catus (Cat).
                                                                                                                                                                                                                                                                                                                                                 AJ409129; CAC41030.1;
                                                                                                                                                                                                                                                                                                                                                                                AJ409130; CAC41033.1;
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Les 2; Conservative
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Best Local Similarity
Matches 2; Conserv
                                                                                                                                                         SEQUENCE FROM N.A.
                                                                                                       NCBI_TaxID=9685;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mitochondrion.
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094PX6 **094PX6**

RESULT 7 Q94PX6

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EMBL; EMBL; EMBL; EMBL; EMBL;

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Query Match

Best Loc Matches

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Q94VA7

RESULT 8 Q94VA7 ID Q94VA

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STRAINENINE MILE Phase I;
MEDLINE=98348442; PubMed=9683477;
Millems H., Jacger C., Baljer G.;
Willems H., Jacger C., Baljer G.;
Willems H., Jacger C., Baljer G.;
J. Bacteriol and genetic map of the obligate intracellular bacterium Coxiella burnetii.";
J. Bacteriol 180:3816-3822(1998).
MEL; AR064963; AAD09947.1;
NON_TER
                                                                                                                             8 AA; 993 MW; 046B5AA453772727 CRC64;
                                                                                                                                                       29.0%; Score 14.5; DB 2; 60.0%; Pred. No. 8.3e+05; iive 1; Mismatches 0;
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SEQUENCE FROM N.A.
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3 WN-GLK 7
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                                                                                                  proliferation of hepatic stellate cells.";
J. Biochem. 127:23-27(2000).
-!- FUNCTION: MAY ACT AS A NEGATIVE EFFECTOR IN THE REGULATION OF THE HEPATIC STELLATE CELLS (ACS). ALSO INHIBITS THE GROWTH OF BOVINE ENDOTHELIAL CELLS AND 316 FIBROBLASTS.
-!- SIMILARITY: IDENTICAL TO THE 63-70 AA REGION OF THE RAT ZAG
 Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Rattus.
                                                                                                                                                                                                                                                                Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                        01-NOV-1996 (TrEMBLrel. 01, Created)
01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
Dutative ORF (Fragment).
Alfalfa mosaic virus.
Viruses; ssRNA positive-strand viruses, no DNA stage; Bromoviridae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE FROM N.A.
MEDLINE=81124289; PubMed=6927843;
Koper-Zwarthoff E.C., Brederode F.T.M., Veeneman G., van Boom J.H.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BOI J.F.;

"Nuclectide sequences at the 5'-termini of the alfalfa mosaic virus
RNAs and the intercistronic junction in RNA 3.";

Nucleic Acids Res. 8:5635-5647(1980).

EMBL; VO0047; CAA23416.1; -

SEQÜENCE 8 AA; 917 MW; 69D40B0775A355B8 CRC64;
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                                                 STRAIN=Sprague-Dawley, TISSUE=Liver,
MEDLINE=20198203, PubMed=10731662,
Kim K.-Y., Choi I., Kim S.-S.,
"Purification and characterization of a novel inhibitor of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Coxiella burnetii.
Bapteria; Froteobacteria; Gammaproteobacteria; Legionellales;
Coxiellaceae; Coxiella.
NCBI_TaxID=777;
                                                                                                                                                                                                                                     30.0%; Score 15; DB 11; Length 8; 100.0%; Pred. No. 8.3e+05;
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01-NOV-1998 (TrEMBLrel. 08, Last sequen
01-NOV-1998 (TrEMBLrel. 19, Last annota
Hypothetical 1.0 kDa protein (Fragment)
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         NCBI_TaxID=10116;
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Eukaryota, Viridiplantae, Streptophyta, Embryophyta, Tracheophyta, Spermatophyta, Magnoliophyta, edicotyledons, core eudicots, Rosidae, eurosids I; Eabales; Fabaceae, Papilionoideae, Phaseoleae, Glycine.

NCBI_TaxID=3847;
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STRAINECV. ESSEX; TISSUE-ROOL;
MEDLINE=91367679; PubMed=1891369;
Laux T., Goldberg R.B.;
"A plant DNA binding protein shares highly conserved sequence motifs
                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                             Xenopus laevis (African clawed frog).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Amphibia; Batrachia; Anura; Mesobatrachia; Pipoidea; Pipidae;
Xenopodinae; Xenopus.
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01-JUN-1998 (TrEMBLrel. 06, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
                           01-JUN-2001 (TrEMBLrel. 17, Created)
01-JUN-2001 (TrEMBLrel. 17, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
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PRT;
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STRAINERF-1;
MEDLINE-99231861; PubMed=10217509;
Huang T.C., Lin R.F., Chu M.K., Chen H.M.;
"Organization and expression of nitrogen-fixation genes in the aerobic nitrogen-fixing unicellular cyanobacterium Synechococcus sp. strain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synechococcus sp. (strain PCC 8801 / RF-1) (Cyanothece PCC 8801). Bacteria; Cyanobacteria; Chrococcales; Cyanothece.
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26.0%; Score 13; DB 2; Length 8;
Best Local Similarity 100.0%; Pred. No. 8.3e+05;
Matches 2; Conservative 0; Mismatches 0; Indels
                                                                                                                                                        STRAIN-CV. ESSEX; TISSUE=ROOL;
Mahalingam R., Knap H.T.;
Submitted (FEB-1998) to the EMBL/GenBank/DDBJ databases.
EMBL; AF047050; AAC03556.1; -.
NON TER 1 1
SEQÜENCE 7 AA; 850 MW; 6AAAAB378637810 CRC64;
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01-JUL-1997 (TrEMBLrel. 04, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
NifH (Fragment)
NIFH.
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BMEL; AF001780; AAC33369.1; -.
NON TER
SEQÜENCE 8 AA, 985 MW; F16B59CDD046C406 CRC64;
with HMG-box proteins.";
Nucleic Acids Res. 19:4769-4769(1991).
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Search completed: November 17, 2003, 18:41:42 Job time : 33 secs

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Sequence 160, Apple Sequence 10, Appl Sequence 10, Appl Sequence 11, Appl Sequence 19, Appl Sequence 19, Appl Sequence 19, Appl Sequence 18, Appl Sequence 18, Appl Sequence 18, Appl Sequence 18, Appl Sequence 28, Appl Sequence 2

Sequence 1, Appli Sequence 16, Appl

Sequence 6, Applisequence 6, Applisequence 14, Applisequence 14, Applisequence 18, Applisequence 20, Applisequence 20, Applisequence 6, Applisequence 14, Applisequence 15, Applisequence 16, Applisequence

Perfect score:

Run on:

Sequence:

Scoring table:

Searched:

Database :

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GENERAL INCREMENTALY

APPLICANT: Barney, S.

APPLICANT: Guthrie, K.

APPLICANT: Guthrie, K.

APPLICANT: Merutka, G.

APPLICANT: Anwer, M.

APPLICANT: Lambert, D.

TITLE OF INVENTION: HYENDEPTIDES WITH ENHANCED PHARMACOKINETIC

TITLE OF INVENTION: PROPERTIES

FILE REFERENCE: 7872-100

CURRENT APPLICATION NUMBER: US/10/351,641

PRIOR APPLICATION NUMBER: 09/350,641

PRIOR FILING DATE: 1999-07-09

PRIOR FILING DATE: 1999-07-09

PRIOR FILING DATE: 1999-05-20

PRIOR FILING DATE: 1999-05-20

PRIOR FILING DATE: 1999-06-20

NUMBER OF SEQ ID NOS: 1757

SOFTWARE: FastSEQ for Windows Version 3.0

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Score 33.5; DB 12;
Pred. No. 5.9e+05;
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Sequence 10, Appl
Sequence 11, Appl
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Sequence 113, Appl
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Sequence 13, Appl
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Sequence 43, Appl
Sequence 4, Appli
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                                                                                                                                                                                                           November 17, 2003, 18:41:46 ; Search time 28 Seconds (without alignments) 52.160 Million cell updates/sec
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(c) 1993 - 2003 Compugen Ltd
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1 US-10-105-293-13
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3 US-10-016-293-11
3 US-10-193-709-17
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US-09-873-676-97
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                                                                                   FEATURE:
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APPLICANT: Zimmermann, Rainer; Park, John E.;
APPLICANT: Zimmermann, Rainer; Park, John E.;
APPLICANT: Rettig, Wolfgang; Old, Lloyd J.
TITLE OF INVENTION: ISOLATED DIMERIC FIBROBLAST ACTIVATION PROTEIN
TITLE OF INVENTION: ISOLATED DIMERIC FIBROBLEST ACTIVATION PROTEIN
TITLE OF INVENTION: ALPHA, AND USES THEREOF
CORRESPONDENCES: 10
CORRESPONDENCES: 10
CORRESPONDENCE: Reife & Lynch
STREET: 805 Third Avenue
CITY: New YORK
CITY: New YORK
COUNTRY: USA
ZIP: 10022
ZIP: 10022
ZIP: 10022
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.5 inch, 2.0 MB storage
COMPUTER: IBM PS/2
OPERATING SYSTEM: PC-DOS
SOFTWARE: Wordperfect
CURRENT APPLICATION DATA:
PILING DATE:
COMPUTER: COMPUTER: DISKETTENTION DATA:
COMPUTER: DATA:
COM
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                Sequence 13, Application US/10016283

Publication No. US20020164702A1

GENERAL INFORMATION:

APPLICANT: Valenzuela et al., David M.

TITLE OF INVERNION: NOVEL TYROSINE KINASE RECEPTORS AND LIGANDS
FILE REFERENCE: REG195-B-PCT-US

CURRENT APPLICATION NUMBER: US/10/016,283

CURRENT PILING DATE: 1998-09-10

PRIOR PELING DATE: 1998-09-10

PRIOR PELING DATE: 1998-09-10

PRIOR PELING DATE: 1996-12-13

NUMBER OF SEQ ID NOS: 36

SEQ ID NO 13

LENGTH: 6
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; OTHER INFORMATION: Description of Artificial Sequence: primer
US-10-016-283-13
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REFERRUCE/DOCKET NUMBER: LUD 5330.1
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 56.0%; Score 28; Best Local Similarity 100.0%; Pred. No. Matches 4; Conservative 0; Mismatch
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PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: US/08/619,280
FILING DATE: 18-MARCH-1996
APPLICATION UNDBER: 08/230,491
FILING DATE: 20-APRIL-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 4, Application US/09265606
Patent No. US20020034789A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEPHONE: (212) 688-9200
TELEFAX: ,(212) 838-3884
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 WSYG 4
US-10-016-283-13
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Publication No. US20030194762A1

| Publication No. US20030194762A1
| GENERAL INFORMATION:
| APPLICANT: KUBOTA, Michio | APPLICANT: HIGASHIYANA, Takanobu | APPLICANT: MIYAKE, Toobio | TITLE OF INVENTION: AAME | TITLE OF INVENTION: SAME | TITLE NETRINGEROUS | T
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ô
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-10-016-283-3

i Sequence 3, Application US/10016283

i Sequence 3, Application US/10016283

publication No US20020164702A1

i GENERAL INFORMATION:

i TITLE OF INVENTION: NOVEL TYROSINE KINASE RECEPTORS AND LIGANDS

i TITLE OF INVENTION: NOVEL TYROSINE KINASE RECEPTORS AND LIGANDS

i TITLE OF INVENTION: NOVEL TYROSINE KINASE RECEPTORS AND LIGANDS

i CURRENT APPLICATION NUMBER: US/10/016,283

i CURRENT PILING DATE: 2001-11-30

i PRIOR APPLICATION NUMBER: PCT/US96/20696

i PRIOR FILING DATE: 1996-12-13

i NUMBER OF SEQ ID NOS: 36

i SOFTWARE: Patentin Ver. 2.0

i ENGTH: 6

i LENGTH: 6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 25; DB 14; Length 6; Pred. No. 5.9e+05; 1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                         The first Xaa is either Trp or Phe
                                                                                                                                                                                                                                                                                                                                                                                                                            56.0%; Score 28; DB 9; Length 7; 100.0%; Pred. No. 5.9e+05; ive 0; Mismatches 0; Indels
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CURRENT FILING DATE: 2002-12-13
PRIOR APPLICATION NUMBER: JP 233364/2000
PRIOR FILING DATE: 2000-08-01
PRIOR FILING DATE: 2000-08-02
PRIOR FILING DATE: 2000-08-02
PRIOR APPLICATION NUMBER: PCT/JP01/06412
PRIOR FILING DATE: 2001-07-25
NUMBER OF SEQ ID NOS: 19
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75.0%;
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ORGANISM: Artificial Sequence
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 7 amino acids
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Matches 3; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             4; Conservative
                                                                                                                                                                                                                                                                                         OTHER INFORMATION:
                                                                                                                                            TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
Matches 4; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 WSYG 4
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APPLICANT: Anwer, M.
APPLICANT: Lambert, D.
TITLE OF INVENTION: HYBRID POLYPEPTIDES WITH ENHANCED PHARMACOKINETIC
TITLE OF INVENTION: PROPERTIES
FILE REFERENCE: 7872-100
CURRENT APPLICATION NUMBER: US/10/351,641
CURRENT FILING DATE: 1999-07-09
PRIOR FILING DATE: 1999-07-09
PRIOR FILING DATE: 1999-07-09
PRIOR FILING DATE: 1999-05-20
PRIOR FILING DATE: 1999-05-20
PRIOR FILING DATE: 1998-05-20
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TITLE OF INVENTION: SYDEMPLIAXIS AND THERAPY OF ACQUIRED IMMUNODEFICIENCY
TITLE OF INVENTION: SYDEMONE
FITLE OF INVENTION: SYDEMONE
FILE REPRENCE: UTSC:267USC1
CURRENT APPLICATION NUMBER: US/09/911,838
CURRENT APPLICATION NUMBER: US/09/911,838
PRIOR APPLICATION NUMBER: 07/834,923
PRIOR FILING DATE: 1992-02-13
PRIOR FILING DATE: 1992-02-13
SOFTWARE: PARCHIN VUMBER: 07/834,923
PRIOR FILING DATE: 1992-02-13
SOFTWARE: PARCHIN VUMBER: 07/834,923
SOFTWARE: PARCHIN VUMBER: 07/834,923
PRIOR FILING DATE: 1992-02-13
SEQ ID NOS: 226
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                                                                                             FEATURE:
) FEATURE:
) OTHER INFORMATION: Description of Artificial Sequence: conserved
) OTHER INFORMATION: motif
US-10-193-709-17
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                                                                                                                                                                                                                                               Length 7;
                                                                                                                                                                                                                                                                                                       0; Indels
                                                                                                                                                                                                                                               Score 24; DB 15;
Pred. No. 5.9e+05;
                                                                                                                                                                                                                                                                                                          Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 919, Application US/10351641 Publication No. US20030186874A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ) OTHER INFORMATION: Core polypeptide US-10-351-641-919
                                     TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                               48.0%;
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                                                                                                                                                                                                                      Query Match
Best Local Similarity 75.0°
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Barney, S. APPLICANT: Guthrie, K. APPLICANT: Merutka, G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
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4 WSFG 7
                                                                                                                                                                                                                                                                                                                                                                     1 WSYG 4
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US-09-911-838-173
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         LENGTH: 7
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Publication No. US20030092079A1

GENERAL INFORMATION:

GENERAL INFORMATION:

APPLICANT: Schlessinger, Joseph

APPLICANT: Schlessinger, Joseph

APPLICANT: Schlessinger, Joseph

APPLICANT: Bendergast, Ann

ITILE OF INVENTION: METHODS FOR IDENTIFYING COMPOUNDS FOR TREATMENT OF CELL

ITILE OF INVENTION: PROLIFERATIVE DISORDERS ASSOCIATED WITH ADAPTOR PROTEIN INTERACT

FILE REFERENCE: 7683-158

CURRENT APPLICATION NUMBER: US/09/393,585

PRIOR FILING DATE: 1995-00-09

PRIOR FILING DATE: 1995-05-24

PRIOR FILING DATE: 1995-05-24

PRIOR FILING DATE: 1995-09-28

PRIOR FILING DATE: 1994-05-20

PRIOR FILING DATE: 1994-05-20
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Sequence 11, Application US/10016283

Publication No. US20020164702A1

GENERAL INFORMATION:

APPLICANT: Valenzuela et al., David M.

TITLE OF INVENTION: NOVEL TYROSINE KINASE RECEPTORS AND LIGANDS

CURRENT APPLICATION NUMBER: US/10/016,283

CURRENT APPLICATION NUMBER: US/20/077,955A

PRIOR PILING DATE: 1998-09-10

PRIOR PILING DATE: 1998-09-10

PRIOR FILING DATE: 1998-12-13

NUMBER OF SEQ ID NOS: 36

SEQ ID NOS: 36

LENGTH: 6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 48.0%; Score 24; DB 14; Length 6; Best Local Similarity 75.0%; Pred. No. 5.9e+05; Matches 3; Conservative 1; Mismatches 0; Indels
                                                                                                                                                                                                                Query Match 50.0%; Score 25; DB 12; Length 8; Best Local Similarity 60.0%; Pred. No. 5.9e+05; Matches 3; Conservative 2; Mismatches 0; Indels
                                                                                                                         ; ORGANISM: Bacillus globisporus
US-10-089-549-10
SOFTWARE: Patentin version 3.1
SEQ ID NO 10
LENGTH: 8
TYPE: PRI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                      1 WSYGL 5
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1 WAFGL 5
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Sequence 97, Application US/09873676
Patent No. US20020077289A1
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Sim, Kim L.
TITLE OF INVENTION: Angiotetatin and Endostatin Binding Proteins and Methods of Use
TILE REFERENCE: 65213-0378 (43170-25933)
CURRENT FILING DATE: 2001-06-04
PRIOR APPLICATION NUMBER: US 60/209,065
PRIOR PLING DATE: 2000-06-02
PRIOR FILING DATE: 2000-06-02
PRIOR FILING DATE: 2001-05-08
NUMBER OF SEQ ID NOS: 123
SOFTWARE: Patentin version 3.1
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                                                                Gaps
                                                                                                                                                                                                                                                                                                                                APPLICANT: May, Michael J
APPLICANT: May, Michael J
APPLICANT: Ghods, Sankar
APPLICANT: Findels, Mark A
APPLICANT: Phillips, Kathryn
APPLICANT: Hannig, Gerhard
TITLE OF INVENTION: ANTI-INFLAMMATORY COMPOUNDS AND USES THEREOF
TITLE REFERENCE: PPI-119
CURRENT APPLICATION NUMBER: US/09/847,946A
CURRENT FILING DATE: 2001-05-02
PRIOR APPLICATION NUMBER: 60/201,261
PRIOR FILING DATE: 2000-05-02
PRIOR PILING DATE: 2000-06-05
PRIOR FILING DATE: 2000-08-22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 OTHER INFORMATION: Description of Artificial Sequence:NBD peptide
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           44.0%; Score 22; DB 10; Length 6; 100.0%; Pred. No. 5.9e+05; tive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                  Sequence 15, Application US/09847946A Publication No. US20030054999A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQ ID NOS: 160
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 15
              Query Match
Best Local Similarity 100.
Matches 3; Conservative
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US-09-873-676-97
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APPLICANT: MacDonald, Nicholas J.
APPLICANT: Sim, Kim L.
TITLE OF INVENTION: Angiostatin and Endostatin Binding Proteins and Methods of Use
TITLE OF INVENTION: Angiostatin and Endostatin Binding Proteins and Methods of Use
FILE REFERENCE: 05213-0378 (43170-259333)
CURRENT FILING DATE: 2001-06-04
PRIOR PILING DATE: 2000-06-02
PRIOR FILING DATE: 2000-06-02
PRIOR FILING DATE: 2001-05-08
NUMBER OF SEQ ID NOS: 123
SOFTWARE: Patentin version 3.1
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; Patent No. US20020156000A1
; Patent No. US20020156000A1
; GENERAL INFORMATION:
    APPLICANT: Ghosh, Sankal J.
    TITLE OF INVENTION: ANTI-INFLAMMATORY COMPOUNDS AND USES THEREOF
; TITLE OF INVENTION: ANTI-INFLAMMATORY COMPOUNDS AND USES THEREOF
; CURRENT APPLICATION NUMBER: US/09/847,940B
; CURRENT PILING DATE: 2001-05-02
; PRIOR FILING DATE: 2000-08-22
; NUMBER OF SEQ ID NOS: 27
; SOFTWARE: Patentin Ver. 2.0
                                                             FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: Synthetic
COTHER INFORMATION: Peptide
US-09-911-838-173
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; OTHER INFORMATION: Description of Artificial Sequence:NBD mutants US-09-847-940B-15
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Pred. No. 5.9e+05;
1; Mismatches 3; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                               ; Sequence 106, Application US/09873676; Patent No. US20020077289A1; GENERAL INFORMATION:
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                                                                                                                                                                             Query Match
Best Local Similarity 42.9%;
Matches 3; Conservative
                  TYPE: PRT ORGANISM: Artificial Sequence
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Best Local Similarity 100.0
Parana 3; Conservative
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1 WDQSLKP 7
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LENGTH: 6
LENGTH: 7
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APPLICANT: Li, Yunger, Stephen D
TITLE OF INVENTION: Expression and Export of Angiogenesis Inhibitors as
TITLE OF INVENTION: Expression and Export of Angiogenesis Inhibitors as
TITLE OF INVENTION: Expression and Export of Angiogenesis Inhibitors as
TITLE OF INVENTION: Expression and Export of Angiogenesis Inhibitors as
TITLE OF INVENTION: Immunofusins
FILER REFERENCE: LEX-006C1
CURRENT FILING DATE: 2002-11-12
PRIOR PELICATION NUMBER: US 60/097,883
PRIOR PELING DATE: 1999-0825
PRIOR PELING DATE: 1999-08-25
NUMBER OF SEQ ID NOS: 54
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 43
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 4, Application US/10239555A
| Sequence 4, Application US/10239555A
| Publication No. US20030186325A1
| GENERAL INFORMATION:
| APPLICANT: BARRY, SIMON
| APPLICANT: HORGAN, CARMEL
| APPLICANT: HORGAN, CARMEL
| TITLE OF INVENTION: METHOD OF SCREENING FOR INHIBITORS OF
| TITLE OF INVENTION: OSTEOPONTIN
| FILE REFERENCE: PG3848
| CURRENT FILING DATE: 2003-03-17
| PRIOR FILING DATE: 2001-03-23
| PRIOR FILING DATE: 2000-03-23
| NUMBER OF SEQ ID NOS: 56
| SEQ ID NO 4
| LENGTH: 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 44.0%; Score 22; DB 12; Length 7; Best Local Similarity 100.0%; Pred. No. 5.9e+05; Matches 3; Conservative 0; Mismatches 0; Indels
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                               Indels
  100.0%; Pred. No. 5.9e+05;
tive 0; Mismatches 0;
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Publication No. US20030139365A1
GENERAL INFORMATION:
APPLICANT: Lo, Kin-Ming
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: PRT ORGANISM: Artificial Sequence
                               3; Conservative
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ORGANISM: Homo sapiens
Best Local Similarity
Matches 3; Conserv
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US-10-292-418-43
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Best Local Similarity 100.0%; Pred. No. 5.9e+05;
Matches 4; Conservative 0; Mismatches 0; Indels 0; Gaps
Qy 3 YGLR 6
Db 4 YGLR 7
Search completed: November 17, 2003, 18:46:56
Job time: 29 secs
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/label= NH2
/note= "amidated glycine"
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OTHER INFORMATION:
OTHER INFORMATION:
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GY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Virginia
: USA
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                                                                                                                                                                                                                                                  RESULT 1
US-08-343-883-2
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Sequence 2, Appli
Sequence 1000, Ap
Sequence 15, Appli
Sequence 15, Appli
Sequence 13, Appli
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                                                                     November 17, 2003, 18:39:46; Search time 21 Seconds (without alignments) 16:118 Million cell updates/sec
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                                                                                                                                                                                                                                                                                                                          /cgn2_6/ptodata/1/iaa/5A_COMB.pep:*
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/cgn2_6/ptodata/1/iaa/6A_COMB.pep:*
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/cgn2_6/ptodata/1/iaa/PcTUS_COMB.pep:*
          GenCore version 5.1.6
(c) 1993 - 2003 Compugen Ltd.
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US-09-082-279B-1000
US-09-315-304B-1000
US-09-834-784-1000
US-08-871-689-3
PCT-US93-07923-15
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US-08-644-271-13

US-08-644-271-13

US-08-619-280A-4

US-08-619-280A-4

US-09-63-947-4

US-08-332-0718-12

US-08-44-271-3

US-08-644-271-3

US-09-644-271-3

US-09-077-955-3

US-08-077-955-3

US-08-286-265-4

US-08-218-837-30
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PCT-US94-01321-63
US-08-522-326-17
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Maximum Match 100%
Listing first 45 summaries
                                                  OM protein - protein search, using sw model
                                                                                                                                                            BLOSUM62
Gapop 10.0 , Gapext 0.5
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                                                                                                             US-09-462-089-3
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Match Length
                                                                                                                                       1 WSYGLRPG 8
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                                                                        Run on:
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APPLICANT: Diffour, Raymond J.
APPLICANT: Diffour, Claude J.M.
APPLICANT: Chouvet, Claire D.
APPLICANT: Chouvet, Claire D.
APPLICANT: Bonneau, Michel B.
TITLE OF INVENTION: Method for improving the organoleptic
TITLE OF INVENTION: qualities of the meat from uncastrated male domestic
TITLE OF INVENTION: animals, vaccines which are usable in this method, new
TITLE OF INVENTION: peptide, in particular for producing these vaccines...
NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
ADDRESSEE: Larson and Taylor
                        Sequence 8, Appli
Sequence 8, Appli
Sequence 17, Appl
Sequence 17, Appl
Sequence 919, App
Sequence 919, App
Sequence 115, Appl
Sequence 2, Appli
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2, Appli
2, Appli
60, Appl
56, Appl
22, Appl
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COMPUTER READABLE FORM:

MEDLUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOSTWARE: PatentIn Release #1.0, Version #1.25

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/343,883

FILING DATE: 17-NOV-1994

CLASSIFICATION: 424

PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 07/946,495 ·

FILING DATE: 09-NOV-1992

PRIOR APPLICATION DATA:

APPLICATION NUMBER: FR 9102513

PRIOR APPLICATION DATA:

APPLICATION NUMBER: FR 9102513

PRIOR APPLICATION DATA:

APPLICATION NUMBER: FR 9115289

FILING DATE: 10-DEC-1991

INFORMATION FOR SEC ID NO: 2:

SEQUENCE CHARACTERISTICS:

LEMNOTH: 8 amino acids
US-09-077-955-11

US-08-472-555-8

US-08-207-575-8

US-08-207-575-8

US-09-393-885-17

US-09-083-798-919

US-09-135-304-919

US-09-135-304-919

US-09-135-304-919

US-09-135-304-919

US-08-125-574-15

US-08-125-574-15

US-08-125-574-15

US-08-125-314-55

US-08-125-314-55

US-08-135-314-163

US-08-175-81-163

US-08-175-81-163

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US-08-175-81-163
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727 Twenty-Third Street, South
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 2, Application US/08343883
Patent No. 5573767
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Sequence 4, Application US/08871689
Patent No. 5955080
GENERAL INFORMATION:
APPLICANT: REILLY, WAYNE G.
APPLICANT: WHITTAKER, ROBERT G.
APPLICANT: FINNEY, KENNETH G.
APPLICANT: FINNEY, KENNETH G.
TITLE OF INVENTION: SELF-ADJUVANTING PEPTIDE VACCINE
TITLE OF INVENTION: DELIVERY SYSTEM AND PRODUCTION THEREOF
NUMBER OF SEQUENCES: 4
                                                                                                                                       88.0%; Score 44; DB 2; Length 7; 100.0%; Pred. No. 2.5e+05; tive 0; Mismatches 0; Indels
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ZIP: 22314

COMPUTER READALLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/871,689
FILING DATE: 09-JUN-1997
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/185,878
FILING DATE: 03-MAY-1994
ATTORNEY/AGENT INFORMATION:
NAME: MILLS, DEMETRA J.
REGISTRATION NUMBER: 1451-004
TELEPHONE: 703-644-1111
TELEPHONE: 703-644-1111
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 40; DB 2;
Pred. No. 2.5e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
ADDRESSEE: LOWE, PRICE, LEBLANC & BECKER STREET: 99 CANAL CENTER PLAZA, SUITE 300 CITY: ALEXANDRIA STATE: VIRGINIA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       80.0%;
nilarity 85.7%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELERY: 703-bo.
TELER: AMERPAT
INFORMATION FOR SEQ ID NO: 4
SEQUENCE CHARACTERISTICS:
SEQUENCE CHARACTERISTICS:
                                                                                                                          Ouery Match
Best Local Similarity 100.
                                TOPOLOGY: linear
MOLECULE TYPE: protein
HYPOTHETICAL: NO
JANTI-SENSE: NO
US-08-871-689-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MOLECULE TYPE: protein HYPOTHETICAL: NO
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      linear
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Best Local Similarity
Matches 6; Consery
                                                                                                                                                                                                                1 WSYGLRP 7
                                                                                                                                                                                                                                        1 WSYGLRP 7
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STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; ANTI-SENSE: NO US-08-871-689-4
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US-08-871-689-4
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APPLICANT: WHITTAKER, ROBERT G.
APPLICANT: WHITTAKER, ROBERT G.
APPLICANT: JENNINGS, PHILLIP A.
APPLICANT: FINNEY, KENNETH G.
TITLE OF INVENTION: SELF-ADJUVANTING PEPTIDE VACCINE
TITLE OF INVENTION: BELF-ADJUVANTING PEPTIDE VACCINE
TITLE OF INVENTION: DELIVERY SYSTEM AND PRODUCTION THEREOF
NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
ADDRESSEE: LOWE, PRICE, LEBLANC & BECKER
STREET: 99 GANAL CENTER PLAZA, SUITE 300
CITY: ALEXANDRIA
STATE: VIRGINIA
COUNTRY: U.S
                                                                                                                                                                                                                                                                                                                                                                                       Query Match 100.0%; Score 50; DB 1; Length 8; Best Local Similarity 100.0%; Pred. No. 2.5e+05; Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Luteinizing hormone-relaesing hormone (LF activity of some synthetic polypeptides. Fragments shorter than decapeptide.
                                                                                                                                                                                                                                                                                                                                         2: FROM 1 TO 8
                                                                                                                                                                                                                                                     Biochem. Biophys. Res. Commun
                                                                                                                                                                                                                                                                                                                                       RELEVANT RESIDUES IN SEQ ID NO:
US-08-343-883-2
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REFERENCE/DOCKET NUMBER: 1451-
TELECHNUTICATION INPORMATION:
TELEPHONE: 703-684-1111
TELEFAX: 703-684-1124
TELERX: AMERPAT
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 7 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 2, Application US/08871689
Patent No. 5955080
                                                                   Redding, T. W. Geiger, R. Konig, W. Wissman, H. Jaeger, G. Sandow, J.
                    Schally, A. V. Arimura, A.
   PUBLICATION INFORMATION:
                                                        Carter, W. H.
                                                                                                                                                                                 Yanaihara,
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                                                                                                                                                                                                                                                                                                                        1972
                                                                                                                                                                                                  TITLE: Lu
TITLE: ac
TITLE: Fr
JOURNAL:
VOLUME: 4
ISSUE: 2
                                                     AUTHORS:
AUTHORS:
AUTHORS:
AUTHORS:
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AUTHORS:
AUTHORS:
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AUTHORS:
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US-08-871-689-2
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DATE:
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US-09-834-784-1000
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US-09-315-304B-1000

i Sequence 1000, Application US/09315304B

i Patent No. 634856B

i GENERAL INFORMATION:

APPLICANT: Barney, S.

APPLICANT: Guthite, K.

APPLICANT: Anwer, M.

APPLICANT: Lambert, D.

TITLE OF INVENTION: HYBRID POLYPEPTIDES WITH ENHANCED PHARMACOKINETIC

TITLE OF INVENTION: PROPERTIES

FILE REFERENCE: 7872-052

CURRENT FILING DATE: 1999-05-20

PRIOR FILING DATE: 1999-05-20

NUMBER OF SEQ ID NOS: 1667

SOFTWARE: FastSEQ for Windows Version 3.0

LEGGTH: B.

LEGGTH: B.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 67.0%; Score 33.5; DB 3; Length 8; 87.5%; Pred. No. 2.5e+05; Live 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Anwer, Mohmed APPLICANT: Lambert, Dennis TITLE OF INVENTION: HYBRID POLYPEPTIDES WITH ENHANCED TITLE OF INVENTION: PHARMACOKINETIC PROPERTIES FILE REFERENCE: 7872-043
CURRENT APPLICATION NUMBER: US/09/082,279B
CURRENT FILIKO DATE: 1998-05-20
NUMBER OF SEQ ID NOS: 1515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FastSEQ for Windows Version 3.0
                                                                                                                         US-09-082-279B-1000
; Sequence 1000, Application US/09082279B
; Patent No. 6258782
; GENERAL INPORMATION:
; APPLICANT: Barney, Shawn
; APPLICANT: Guthrie, Kelly
; APPLICANT: Merutka, Gene
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ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity 87.5
Matches 7; Conservative
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1 WSYGLQP 7
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SEQ ID NO 1000
LENGTH: 8
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APPLICANT: REILLY, WAYNE G.
APPLICANT: WHITTAKER, ROBERT G.
APPLICANT: JENNINGS, PHILLIP A.
APPLICANT: JENNINGS, PHILLIP A.
APPLICANT: FINNEY, KENNETH G.
TITLE OF INVENTION: SELF-ADJUVANTING PEPTIDE VACCINE
TITLE OF INVENTION: DELIVERY SYSTEM AND PRODUCTION THEREOF
NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
ADDRESSEE: LOWE, PRICE, LEBLANC & BECKER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Indels
Sequence 1000, Application US/09834784

Sequence 1000, Application US/09834784

GENERAL INFORMATION:
FAPPLICANT: Barney, Shawn

APPLICANT: Guthrie, Kelly
APPLICANT: Anwer, Mohmed
APPLICANT: Lambert, Dennis
TITLE OF INVENTION: PHARMACOKINETIC PROPERTIES
FILE REFERENCE: 7872-043

CURRENT APPLICATION NUMBER: US/09/834,784

CURRENT FILING DATE: 2001-04-13

PRIOR APPLICATION NUMBER: 09/082,279

PRIOR APPLICATION NUMBER: 09/082,279

PRIOR APPLICATION NUMBER: 09/082,279

PRIOR PRING DATE: 1998-05-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPTITE: IBM PC compatible
SOFTWARE: Patentin Release #1.0, Version #1.25
CURSTRIATION DATA:
APPLICATION NUMBER: US/08/871,689
FILING DATE: 09-JUN-1997
CLASSIFICATION: 424
PRICA APPLICATION ADATA:
APPLICATION NUMBER: US 08/185,878
FILING DATE: 03-MAY-1994
FILING DATE: 03-MAY-1994
ATTORNEY/AGENT INCRMATION:
NAME: MILLS, DEMETRA J.
REGISTRATION NUMBER: 34.506
REFERENCE/POCKET NUMBER: 1451-004
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
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Pred. No. 2.5e+05;
0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQ ID NOS: 1515
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 1000
LENGTH: 8
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Patent No. 5955080
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: PRT ORGANISM: Artificial Sequence
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87.5%;
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Best Local Similarity 87.5
7; Conservative
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US-08-644-271-13
Sequence 13, Application US/08644271
Sequence 13, Application US/08644271
Sequence 13, Application US/08644271
Sequence 13, Application US/08644271
TITLE OF INVENTION: NOVEL TYROSINE KINASE RECEPTORS
TITLE OF INVENTION: AND LIGANDS
NUMBER OF SEQUENCES: 32
CORRESPONDENCE ADDRESS:
ADDRESSED: Regeneron Pharmaceuticals, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY: NEW TOTA

COUNTRY: USA

ZIP: 10591

COMPUTER READBALE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/374,834
FILING DATE: 19-JAN-1995
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/095,658
FILING DATE: 21-JUL-1993
ATTORNEY/AGENT INFORMATION:
ANAMERICATION PROPERTY OF ANAMERICATION NUMBER: DEPOSATE TOTAL TOTAL NUMBER: DEPOSATE T
                                                                                                                                                                                                                                          GENERAL INFORMATION:
APPLICANT: Valenzuela, et al.
TITLE OF INVENTION: NOVEL TYROSINE KINASE RECEPTOR
NUMBER OF SEQUENCES: 17
CORRESPONDENCE ADDRESS:
ADDRESSEE: Regeneron Pharmaceuticals, Inc.
STREET: 777 Old Saw Mill River Road
CITY: Tarrycown
STATE: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                56.0%; Score 28; DB 1; Le
100.0%; Pred. No. 2.5e+05;
ative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Cobert, Robert J.
REGISTRATION NUMBER: 36,108
REPERENCE, DOCKET NUMBER: REG 190A
TELECOMMUNICATION INFORMATION:
TELEPHONE: (914) 345-7400
                                                                                                                          RESULT 9
US-08-374-834-13
; Sequence 13, Application US/08374834
; Patent No. 5656473
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STREET: 777 Old Saw Mill Road CITY: Tarrytown STATE: NY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEFAX: (914) 345-7721 INFORMATION FOR SEQ ID NO: 13:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS:
LENGTH: 6 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity 100.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; TOPOLOGY: unknown
; MOLECULE TYPE: peptide
US-08-374-834-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ZIP: 10591
COMPUTER READABLE FORM:
WSYG 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 WSYG 4
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                                   셤
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Ouery Match 56.0%; Score 28; DB 5; Length 5; Best Local Similarity 100.0%; Pred. No. 2.5e+05; Matches 4; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 15, Application PC/TUS9307923
GENERAL INFORMATION:
APPLICANT: Schlossman, Stuart F.
APPLICANT: Schlossman, Stuart F.
APPLICANT: Schlossman, Stuart F.
APPLICANT: Tanaka, Toshiaki
TITLE OF INVENTION: HUMAN CD26 AND METHODS FOR USE
NUMBER OF SEQUENCES: 16
CORRESPONDENCES: 16
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson
STRIE: Massachusetts
COUNTY: Boston
STATE: Massachusetts
COUNTY: U.S.A.
IP: 02110-2804
COMPUTER: IBM PS/2 Model 502 or 553X
COMPUTER: IBM PS/2 Model 502 or 553X
COMPUTER: IBM PS/2 Model 502
COMPUTER: 19930819
CURRENT APPLICATION DATA:
APPLICATION NUMBER: 07/934,162
FILING DATE: 19930819
CLASSIFICATION NUMBER: 07/932,211
FILING DATE: 01-992
APPLICATION NUMBER: 07/932,211
FILING DATE: 01-992
APPLICATION NUMBER: 07/932,211
FILING DATE: 06-FEB-1992
ATTORNEY AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: Fraser, Janis K.
REGISTRATION NUMBER: 34,819
REGISTRATION NUMBER: 34,819
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 542-5070
TELEFAX: 200154
TELEX: 200154
INFORMATION FOR SEQ ID NO: 15:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                     62.0%;
                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity 71...
5. Conservative
                                                                                                                          LENGTH: 7 amino acide
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: protein
HYPOTHETICAL: NO
                            TELEX: AMERPAT
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
      703-684-1124
                                                                                                                             7 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: amino acid
STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 WSYGWLP 7
                                                                                                                                                                                                                                                                                                              ; ANTI-SENSE: NO
US-08-871-689-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 WSYG 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 8
PCT-US93-07923-15
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         TELEFAX:
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Gaps

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Gaps
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Patent No. 5767242
GENERAL INFORMATION:
APPLICANT: Entering, Wolfgang; Old, Lloyd J.
TITLE OF INVENTION: ISOLATED DIMERIC FIBROBLAST ACTIVATION PROTEIN
TITLE OF INVENTION: ALPHA, AND USES THEREOF
NUMBER OF SEQUENCES: 10
CORRESPONDENCE ADDRESS:
ADDRESSEE: Felfe & Lynch
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
APPLICANT: Valenzuela et al., David M.
APPLICANT: Valenzuela et al., David M.
TITLE OF INVENTION: NOVEL TYROSINE KINASE RECEPTORS AND LIGANDS
FILE REFERENCE: REG195-9-EPT-US
CURRENT APPLICATION NUMBER: US/09/07, 955A
CURRENT FILING DATE: 1998-09-10
FEARLIER APPLICATION NUMBER: 08/644, 271
EARLIER FILING DATE: 1996-12-13
EARLIER FILING DATE: 1996-05-10
EARLIER FILING DATE: 1996-05-10
EARLIER FILING DATE: 1996-05-10
SARLIER FILING DATE: 1996-05-10
SARLIER FILING DATE: 1995-12-15
NUMBER OF SEQ ID NOS: 36
SOCTHARE: Patentin Ver. 2.0
SEQ ID NO 13
LENGTH: 6
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; OTHER INFORMATION: Description of Artificial Sequence: primer
US-09-077-955-13
                                                                                                                                                                                                                                                                                                    56.0%; Score 28; DB 2; Length 6; 100.0%; Pred. No. 2.5e+05; Live 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    56.0%; Score 28; DB 4; Length 6; 100.0%; Pred. No. 2.5e+05; ative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 13, Application US/09077955A Patent No. 6413740
    PELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: PRT ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SSEE: Felfe & Lynch
1: 805 Third Avenue
New York City
                          TELEPHONE: 914-345-7400
TELEFAX: 914-345-7721
                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity luv...
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity 100.
                                                                                                                                                                                    unknown
                                                                                       INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 6 amino acids
                                                                                                                                                                                                                               , MOLECULE TYPE: peptide US-08-469-537A-95
                                                                                                                                                                   TYPE: amino acid
STRANDEDNESS: sir
                                                                                                                                                                                                                                                                                                                                                                                                     1 WSYG 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                 3 WSYG 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3 WSYG 6
                                                                                                                                                                                                                 TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 12
US-09-077-955-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-08-619-280A-4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                56.0%; Score 28; DB 2; Length 6; 100.0%; Pred. No. 2.5e+05; Live 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 11
US-08-469-537A-95
15 Sequence 95, Application US/08469537A
15 Patent No. 5843749
16 GENERAL INPORMATION:
17 ILLE OF INVENTION: BHK AND ROR TYROSINE
17 TILLE OF INVENTION: KINASES
NUMBER OF SEQUENCES: 107
17 CORRESPONDENCE ADDRESS:
ADDRESSEE: Regeneron Pharmaceuticals, Inc.
5 TATE 77 Old Saw Mill River Road
CITY: Tarrytown
STATE: NY
COUNTRY: U.S.A.
2IP: 10591
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: Diskette
COMPUTER: Enactor Version 2.0
SOFTWARE: Diskette
COMPUTER: U.S.A.
SOFTWARE: PastERO Version 2.0
CURRENT APPLICATION NUMBER: USSN 08/469,537A
FILING DATE: 17-MAR-1995
APPLICATION NUMBER: USSN 08/144,992
FILING DATE: 28-OCT-1993
APPLICATION NUMBER: USSN 08/144,992
PILLING DATE: 28-OCT-1993
APPLICATION NUMBER: USSN 08/144,992
                                         OPERATING SYSTEM: DOS
OSCTWARE: FASTEM: DOS
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/644,271
FILING DATE: 10-MAY-1996
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: USSN 60/008,657
FILING DATE: 15-DEC-1995
ATTORNEY/AGNT INFORMATION:
NAME: CODERT, ROBERT J
REGISTRATION NUMBER: 36,108
RECISTRATION NUMBER: 36,108
REFERENCE/DOCKET NUMBER: RG 195A
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE: 28-OCT-1993
APPLICATION WINBER: USSN 07/736,559
FILING DATE: 26-UUL-1991
ATTORNEY/AGENT INFORMATION:
NAME: Kempler, Ph. D., Gail M
REFISTRATION NUMBER: 32,143
REFERENCE/DOCKET NUMBER: REG 070C
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: 914-345-7721
TELEX:
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 56.0
Best Local Similarity 100.
Matches 4; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 6 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY: linear
MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-08-644-271-13
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GENERAL INFORMATION:
APPLICANT: ROBERN, ERICH G.
APPLICANT: ROBERN, ERICH G.
APPLICANT: RESSLER, HORST
APPLICANT: RESSLER, HORST
APPLICANT: RENEND, MICHAEL
APPLICANT: RENEND, MICHAEL
APPLICANT: RENEND, MICHAEL
APPLICANT: RICHAEL
APPLICANT: APPLICANT
TITLE OF INVENTION: THEIR DERIVATIVES FOR INCORPORATION IN PHARMACOLOGICALLY
TITLE OF INVENTION: ACTIVE PEPTIDES AND THEIR SALTS
NUMBER OF SEQUENCES: 17
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    51.0%; Score 25.5; DB 1; 75.0%; Pred. No. 2.5e+05;
                                                                                                                                                                                                                                                                                                                                                                                                STREET: 1100 NEW YORK AVENUE, N.W.
CITY: WASHINGTON
COUNTRY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          326/216933
                                                                                                                                                         ; Sequence 12, Application US/08332071B
; Patent No. 5556836
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ATTORNEY/AGENT INFORMATION:
NAME: CHAPIN, MARLANA K.
REGIESTRATION NUMBER: 35,843
REFERENCE/DOCKET NUMBER: 326/
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-861-3000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity 75.v.
6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: 202-822-0944
TELEX: 6714627 CUSH
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 7 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 WS-XLRPG 7
                                          1 SYXLRP 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STRANDEDNESS:
    SYGLRP
                                                                                                              RESULT 15
US-08-332-071B-12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME/KEY: misc_feature
OTHER INFORMATION: Description of Artificial Sequence: synthetic peptide
NAME/KEY: misc_feature
LOCATION: (3)...(3)
OTHER INFORMATION: Xaa is D-Leu, D-Ser(But), D-Trp, (2-napthyl)-D-Ala, or Gly
NAME/KEY: misc_feature
LOCATION: (7)...(7)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Gaps
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Fatent No. 6448031

GENERAL INFORMATION:
APPLICANT: KASHIMOTO, KAZUHISA
APPLICANT: NAGANO, YUMIKO
TITLE OF INVENTION: METHOD FOR PRODUCING LH-RH DERIVATIVES
FILE REFERENCE: 7339-0002-0PCT
CURRENT APPLICATION NUMBER: US/09/463,947

CURRENT FILING DATE: 1997-08-04

PRIOR APPLICATION NUMBER: PCT/JP97/02705

PRIOR APPLICATION NUMBER: PCT/JP97/02705

PRIOR PILING DATE: 1997-08-04

NUMBER OF SEQ ID NOS: 4

SEQ ID NO 4

LENGTH: 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 52.0%; Score 26; DB 4; Length 7; Best Local Similarity 83.3%; Pred. No. 2.5e+05; Matches 3.5; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FEATURE:
OTHER INFORMATION: The first Xaa is either Trp or Phe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 56.0%; Score 28; DB 1; Length 7; Best Local Similarity 100.0%; Pred. No. 2.5e+05; Matches 4; Conservative 0; Mismatches 0; Indels
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.5 inch, 72.0 MB storage COMPUTER: IBM PS/2
OPERATING SYSTEM: PC-DOS
SOFTWARE: Wordperfect
CURRENT APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US/08/619,280A
FILING DATE: 18-MARCH-1996
CLASSIPICATION: 435
PRIOR APPLICATION: 435
APPLICATION NUMBER: 08/230,491
FILING DATE: 20-APRIL-1994
ATTORNEY/AGENT INFORMATION:
NAME: Hanson, No. 5767242man D.
REGISTRATION NUMBER: 10D 5330.1
FELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEPAX: (212) 888-3884
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 7 amino acids
TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 WSYG 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-08-619-280A-4
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GenCore version 5.1.6
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OM protein - protein search, using sw model

November 17, 2003, 18:27:39; Search time 13.5 Seconds (without alignments) 64:112 Million cell updates/sec Run on:

US-09-462-089-4 48 1 GSGSGLRPG 9 Title: Perfect score: Sequence:

Scoring table:

BLOSUM62 Gapop 10.0 , Gapext 0.5

283308 segs, 96168682 residues Searched:

789 Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 9

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

PIR_76:*
1: pirl:*
2: pir2:*
3: pir3:*
4: pir4:* Database :

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES	Description	37 D-amino-acid oxida	30 Iq heavy chain CRD		T-cell	T-cell	3 T	T-cell	carnoci	H	7-1		T-cel	T-cell	T-cell	Ig heav		18 T-cell receptor be	synapt	H	T-cell		peptidylglycine	leucokinin'v - Mad	7.3		H 6I		10.4 411
ns	ID	539437	PT0280	PT0725	C53284	PT0689	PT054	PT0581	A58718	PT0268	PT0572	PT0605	PT0668		PT0719		D58503	PH0918	D44823				A38671	JS0315					15112
	DB	. 2				2	~	~											~								~		
	Length	6	φ	80	2	7	7	7	7	6	S	9	9	9	7	80	6	σ	S	S	9	9	7	80	σ	6	6	'n	4
de	Query Match	43.8	41.7	41.7	39.6	39.6	37.5	37.5	37.5	37.5	35.4	35.4		35.4		35.4	35.4	35.4	33.3	33.3	33.3	33.3	33.3	33.3	33.3	33.3	•	31.2	,
	Score	21	20	20	19	19	18	18	18	18	17	17	17	17	17	17	17	17	16	16	16	16	16	16	16	16	16	15	ב
	Result No.	-	7	m	4	Ω.	9	7	æ	6	10	11	12	13	14	15	16	17	18	19		21	22	23	24		56		ac

244K exoantigen -	T-cell receptor be	T-cell receptor be	peptidyl-dipeptida	spinal cord peptid	copper resistance	T-cell receptor be	T-cell receptor be	T-cell receptor be	T-cell receptor be	Ig H chain V-D-J r	T-cell receptor be	T-cell receptor be	glutathione transf	Ig heavy chain CRD	zymogen granule me
A33098	PT0515	PT0663	JN0862	C23751	A41225	PT0589	PT0593	PT0529	PT0623	PH1618	PT0627	PT0631	S59902	PT0288	PC2197
8	N	7	~	N	~	(7	7	0	~	7	N	~	~	~	0
7	7	7	Ŋ	Ŋ	Ŋ	9	9	7	7	œ	ω	œ	σ	σ	0
31.2	31.2	31.2	29.2	29.2	29.2	29.5	29.2	29.2	29.2	29.2	29.2	29.2	29.2	29.5	29.2
15	15	15	14	14	14	14	14	14	14	14	14	14	14	14	14
30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45

ALIGNMENTS

	RESULT 1 S3947 D-amino-acid oxidase (EC 1.4.3.3) - Trigonopsis variabilis (fragment) C.Species: Trigonopsis variabilis C.Species: Trigonopsis variabilis C.Date: 19-Mar-1997 #sequence_revision 05-Dec-1997 #text_change 07-May-1999 C.Accession: S39437 R.Schraeder, T.; Andreesen, J.R. Eur. J. Biochem. 218, 735-744, 1993 A;Title: Evidence for the functional importance of Cys298 in D-amino acid oxidase from 1A, Reference number: S39437; MUID:94094869; PMID:7903639 A;Accession: S39437 A;Molecule type: protein
	A;Residues: 1-9 <\$CH> A;Experimental source: CBS 4095 C;Fuction: oxidoreductase; catalyzes the oxidation of D-amino acids to their correst, A;Description: oxidoreductase; catalyzes the oxidation of the enzyme by molecular oxygen is accompanied by the release of l C;Keywords: FAD; oxidoreductase
·····	Query Match 43.8%; Score 21; DB 2; Length 9; Best Local Similarity 80.0%; Pred. No. 2.8e+05; Matches 4; Conservative 0; Mismatches 1; Indels 0; Gaps 0;
	Oy 5 GLRPG 9
	RESULT 2 PT0280 Ig heavy chain CRD3 region (clone 4-91B) - human (fragment) C;Species: Homo sapiens (man) C;Species: Homo sapiens (man) C;Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change 16-Aug-1996 C;Accession: PT0280 R;Yamada, M; Wasserman, R.; Reichard, B.A.; Shane, S.; Caton, A.J.; Rovera, G. J. Exp. Med. 173, 395-407, 1991108337; PMID:1899102 A;Title: Preferential utilization of specific immunoglobulin heavy chain diversity and JA;Reference number: PT0222; MUID:91108337; PMID:1899102 A;Accession: PT0280 A;Molecule type: DNA A;Residues: 1-6 <am>A;Residues: 1-6 <amaa;residues: 1-6="" <amaaa;residues:="" <amaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaa<="" td=""></amaa;residues:></am></am></am></am></am></am></am></am></am></am></am></am></am></am>
	O; Mismatch

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carnocin U149 - Carnobacterium sp. (fragment)
C;Species: Carnobacterium sp.
C;Species: 23-dan-1998 #sequence_revision 30-Jan-1998 #text_change 30-Jan-1998
C;Apate: 23-dan-1998 #sequence_revision 30-Jan-1998 #text_change 30-Jan-1998
C;Accession. A58718
R;Stoffels, G; Nissen-Meyer, J; Gudmundsdottir, A; Sletten, K; Holo, H; Nes, I.F.
Appl. Environ. Microbiol. 58, 1417-1422, 1992
Aprille: Purification and characterization of a new bacteriocin isolated from a Carnobac A;Reference number: A58718; MUID:92321768; PMID:1622206
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C;Species: Mus musculus (house mouse)
C;Species: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 30-May-1997
C;Accession: PTO581
R;Feeney, A.J.
B;Feeney, A.J.
J;Title: Junctional sequences of fetal T cell receptor beta chains have few N regions. A;Reference number: PTO509; MUID:91277601; PMID:1711558
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       J. Exp. Med. 174, 115-124, 1991
A,Title: Junctional sequences of fetal T cell receptor beta chains have few N regions.
A,Reference number: PT0509; MUID:91277601; PMID:1711558
A,Accession: PT0543
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C;Date: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 30-May-1997
C;Accession: PT0543
       Gaps
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Pred, No. 2.88+05;
2; Indels
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Pred. No. 2.8e+05;
0; Mismatches 2; Indels
Indels
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A;Residues: 1-7 <FEE>
A;Experimental source: day 18 fetal thymus, strain BALB/c
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A,Experimental source: day 19 fetal thymus, strain BALB/c
C,Keywords: T-cell receptor
       <u>..</u>
       Mismatches
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Best Local Similarity 66.7%;
Matches 4; Conservative
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A,Status: translation not shown
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       3; Conservative
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A, Molecule type: protein
A, Residues: 1-7 <STO>
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       Matches
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PT0581
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C.Species: Oryctolagus cuniculus (domestic rabbit)
C.Species: Oryctolagus cuniculus (domestic rabbit)
C.Species: Oryctolagus cuniculus (domestic rabbit)
C.Saccesion: C53284
R.Harindranath, N.; Alexander, C.B.; Mage, R.G.
Mol. Immunol. 28, 881-888, 1991
A.Title: Evolutionarily conserved organization and sequences of germline diversity and J.A.Accesion: C53284
A.A.Accesion: C53284
A.A.A.Cession: C53284
A.A.A.Cession: C53284
A.A.Cession: C53284
A.A.C
                                                                                                                                               C; Species: Mus musculus (house mouse)
C; Date: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 30-May-1997
C; Date: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 30-May-1997
C; Date: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 30-May-1997
C; Accession: PT0725
A; Pitle: Junctional sequences of fetal T cell receptor beta chains have few N regions.
A; Reference number: PT0509; MUID:91277601; PMID:1711558
A; Accession: PT0725
A; Accession: PT0725
A; Atatus: translation not shown
A; Molecule type: DNA
A; Molecule type: DNA
A; Residues: 1-8 < FEB>
A; Experimental source: newborn thymus, strain BALB/c
C; Keywords: T-cell receptor
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A;Title: Junctional sequences of fetal T cell receptor beta chains have few N regions.
A;Reference number: PT0509; MUID:91277601; PMID:1711558
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C;Species: Mus musculus (house mouse)
C;Date: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 30-May-1997
C;Accession: PT0689
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C, Keywords: T-cell receptor
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41.7%; Score 20; DB 2; I
Best Local Similarity 80.0%; Pred. No. 2.8e+05;
Matches 4; Conservative 0; Mismatches 1;
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A;Status: translation not shown
A;Molecule type: DNA
A;Residues: 1-7 <FEE>
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Best Local Sjmilarity
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GTGGG 5
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J. Exp. Med. 174, 115-124, 1991
A;Tille: Junctional sequences of fetal T cell receptor beta chains have few N regions. A;Rille: Junctional sequences of fetal T cell receptor beta chains have few N regions. A;Recession: PT0668
A;Status: translation not shown
A;Molecule type: mRNA
A;Residues: 1-6 cFEEA
A;Experimental source: day 4 postnatal thymus, strain BALB/c
C;Keywords: T-cell receptor
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J. Exp. Med. 174, 115-124, 1991
A;Title: Junctional sequences of fetal T cell receptor beta chains have few N regions.
A;Reference number: PT0509; MUID:91277601; PMID:1711558
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C;Species: Mus musculus (house mouse)
C;Date: 17-101-1992 #sequence_revision 17-Jul-1992 #text_change 30-May-1997
C;Accession: PT0668
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C;Species: Mus musculus (house mouse)
C;Date: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 30-May-1997
C;Accession: PT0723
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Pred. No. 2.8e+05;
1; Mismatches 0; Indels
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75.0%; Pred. No. 2.8e+05;
M.comprohes 0;
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A;Molecule type: DNA
A;Residues: 1-6 <FEEA
A;Experimental source: newborn thymus, strain BALB/C
C;Keywords: T-cell receptor
A;Status: translation not shown
A;Molecule type: mRNA
A;Residues: 1-6 <FEE
A;Experimental source: newborn thymus, strain BALB/c
C;Keywords: T-cell receptor
                                                                                                                                                          Query Match 35.4%;
Best Local Similarity 75.0%;
Matches 3; Conservative
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Best Local Similarity 75.01
Matches 3; Conservative
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Best Local Similarity
Matches 3; Conserv
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2 SGAG 5
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SGTG 5
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PT0719
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PT0723
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                                                                                                                                                                                                                                                                                                                                     Ig heavy chain CRD3 region (clone 3-94B) - human (fragment)
C; Species: Homo sapiens (man)
C; Species: Homo sapiens
C; Accession: PT0268
R; Yamada, M: Wasserman, R: Reichard, B.A.; Shane, S.; Caton, A.J.; Rovera, G.
J. Exp. Med. 173, 395-407, 1991
A; Reference number: PT0222; MUID:91108337; PMID:1899102
A; Reference number: PT0222; MUID:91108337; PMID:1899102
A; Molecule type: DNA
A; Molecule type: DNA
A; Molecule type: DNA
A; Residues: 1-9 < YAM>A; Residues: 1-9 < YAM>A; Residues: 1-9 < YAM>A; Residues: heterotetramer; immunoglobulin
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A.Title: Junctional sequences of fetal T cell receptor beta chains have few N regions. A.Title: Junctional sequences of fetal T cell receptor beta chains have few N regions. A.Reference number: PT0509; MUID:91277601; PMID:1711558
A.Accession: PT0572
A.Status: translation not shown
A.Status: translation not shown
A.Residues: 1-5 <FEE>
A.Residues: 1-5 <FEE>
A.Residues: 1-5 <FEE>
A.Resperimental source: day 19 fetal thymus, strain BALB/c
C.Keywords: T-cell receptor
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C;Species: Mus musculus (house mouse)
C;Date: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 30-May-1997 C;Accession: PT0605
R;Feeney, A.J.
J. Exp. Med. 174, 115-124, 1991
A;Title: Junctional sequences of fetal T cell receptor beta chains have few N regions. A;Reference number: PT0609; MUID:91277601; PMID:1711558
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                                                       Length 7;
                                                 Score 18; DB 2; I
Pred. No. 2.8e+05;
2; Mismatches 1;
  C; Keywords: antibiotic; lanthionine
                                                 Query Match
Best Local Similarity 50.0%;
Matches 3; Conservative
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1 GSEIQP 6
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ig heavy chain CRD3 region (clone 4-91A) - human (fragment)
C;Species: Homo sapiens (man)
C;Species: Jo.Sep-1993 #text_change 16-Aug-1996
C;Accession: PT0279
R;Yamada, M.; Wasserman, R.; Reichard, B.A.; Shane, S.; Caton, A.J.; Rovera, G.
J. Exp. Med. 173, 395-407, 1991
A;Title: Preferential utilization of specific immunoglobulin heavy chain diversity and A;Accession: PT0279
A;Accession: PT0279
A;Accession: PT0279
A;Molecule type: DNA
A;Molecule type: DNA
A;Residues: 1-8 < YAM>
A;Residues: 1-8 < YAM>
A;Residues: 1-8 < YAM>
A;Experimental source: B lymphocyte
C;Keywords: heterotetramer; immunoglobulin
R;Feeney, A.J.
J. Exp. Med. 174, 115-124, 1991
A;Title: Junctional sequences of fetal T cell receptor beta chains have few N regions.
A;Reference number: PT0509; MUID:91277601; PMID:1711558
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A; Status: translation not shown
A; Residues: 1-7 <FRES
A; Experimental source: newborn thymus, strain BALB/c (clone 140-2F)
A; Accession: PT0538
A; Accession: PT0538
A; Status: translation not shown
A; Molecule type: mRNA
A; Residues: 1-7 <FRES
A; Experimental source: newborn thymus, strain BALB/c (clone 111-1N)
C; Keywords: T-cell receptor
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Cowden C., Stretton A.O.W.;
"Eight novel FMRFamide-like neuropeptides isolated from the nematode
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Peptides 16:491-500(1995).
-!- SIMILARITY: BELONGS TO THE FARP (FMRFAMIDE RELATED PEPTIDE)
FAMILY.
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MEDLINE=95380362; PubMed=7651904;
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                                                                     November 17, 2003, 18:24:14; Search time 10 Seconds (without alignments) 42.324 Million cell updates/sec
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           GenCore version 5.1.6
(c) 1993 - 2003 Compugen Ltd.
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                                                                                                                                                                                        127863 segs, 47026705 residues
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Maximum Match 100%
Listing first 45 summaries
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Match
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Perfect score:
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periplaneta
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Submitted (APR-1999) to the SWISS-PROT data bank.
-!- MISCELLANEOUS: THERE ARE SEVEN EXTERNAL GLYCOPROTEINS IN HSV1 AND
2: GH, GB, GC, GG, GJ, GI, AND GE.
-!- MISCELLANEOUS: GLYCOPROTEIN G IS MUCH LARGER IN HSV-2 THAN IN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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Eukaryota; Metazoa; Nematoda; Chromadorea; Ascaridida; Ascaridoidea;
Ascarididae; Ascaris.
NOBI_TaxID=6253;
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P19149
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Viruses; dSDNA viruses, no RNA stage; Herpesviridae;
Alphaherpesvirinae; Simplexvirus.
NCBI_TaxID=103921;
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15-JUL-1999 (Rel. 38, Last sequence update)
15-JUL-1999 (Rel. 38, Last annotation update)
Glycoprotein G (Fragment)
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01-NOV-1995 (Rel. 32, Last sequence update)
01-FEB-1996 (Rel. 33, Last annotation update)
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               RB42_LITRU
XY1A_STRSQ
XY1A_STRSQ
OCPI_OCTMI
OCPI_OCTMI
ALL4_CARMA
UXA4_CHLTR
CIPI_MYTED
ALL2_CARMA
ALL4_CARMA
ALL4_CARMA
                                                                                                                                                                                                                                                                                                                                                       PRT;
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5; Conservative
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AMIDATION

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35.4%;
75.0%;
                                                                               3; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Oryctolagus cuniculus
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                       Query Match
Best Local Similarity
Matches 3; Conserv
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21-JUL-1986 (
16-OCT-2001 (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DSIP_RABIT
P01158;
                                                                                                                                                                                                                                                                                                                                           LEUMA
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SEQUENCE
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                                                                                                                                                                                                                                                                                   KESULT 5
LCKS_LEUMA
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DSIP_RABI
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    a Carnobacterium sp.";
Appl. Environ. Microbiol. 58:1417-1422(1992).
-!- FUNCTION: LANTHONIBE-CONTAINING PEPTIDE ANTIBIOTIC (LANTIBIOTIC)
ACTIVE ON GRAM-POSITIVE BACTERIA.
Antibiotic; Bacteriocin; Lantibiotic.
                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDLINE=92321768; PubMed=1622206;
Stoffels G., Nissen-Meyer J., Gudmundsdottir A., Sletten K., Holo H.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         "Purification and characterization of a new bacteriocin isolated from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             "Plasma protein map: an update by microsequencing.";
Electrophoresis 13:707-714(1992).
-!- MISCELLANBOUGS: ON THE 2D-GEL THE DETERMINED PI OF THIS UNKNOWN
PROTEIN IS: 5, ITS WM IS: 48 kDa.
SWISS-2DPAGE; P30092; HUMAN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
NCBI_TaxID=9606;
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01-APR-1993 (Rel. 25, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
Unknown protein from 2D-page of plasma (Spot 14) (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Carnobacterium sp. (strain U149).
Bacteria, Firmicutes, Lactobacillales, Carnobacteriaceae,
Carnobacterium.
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Hughes G.J., Frutiger S., Paquet N., Ravier F., Pasquali
Sanchez J.-C., James R., Tissot J.-D., Bjellqvist B.,
Hochstrasser D.F.;
                                                        Score 19; DB 1; Length 9;
Pred. No. 1.3e+05;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               37.5%; Score 18; DB 1; Length 7; 50.0%; Pred. No. 1.3e+05; cive 2; Mismatches 1; Indels
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524F073774176877 CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                01-JUN-1994 (Rel. 29, Created)
01-JUN-1994 (Rel. 29, Last sequence update)
116-QCT-2001 (Rel. 40, Last annotation update)
Lancibiotic carnocin U149 (Fragment).
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                                                  h 39.6%;
Similarity 66.7%;
4; Conservative
     1012 MW;
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Best Local Similarity 50...
3; Conservative
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                                                  Query Match
Best Local Similarity
Matches 4; Conserv
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GSEIQP 6
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     9 . A. .
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UPA6_HUMAN
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LANC_CARUI
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MEDLINE=87052651; PubMed=2877794;
MEDLINE=87052651; PubMed=2877794;
MEDLINE=87052651; PubMed=2877794;
MEDLINE=87052651; PubMed=2877794;
MEDLINE=87052651; PubMed=287794;
MISOLation, primary Structure, and synthesis of leucokinins V and VI:
myotropic peptides of Leucophaea maderae.";
Comp. Biochem. Physiol. 88C:27-30(1987).
-!- FUNCTION: THIS CEPHALONYOTROPIC PEPTIDE STIMULATES CONTRACTILE
ACTIVITY OF COCKROACH PROTODEUM (HINDGUT).
-!- STMILARITY: TO THE OTHER LEUCOKININS.
-!- PINTIARIARITY: TO THE OTHER LEUCOKININS.
Meuropeptide; Amidation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE, AND SYNTHESIS.
MEDLINE=7905441; PubMed=568769;
Schoenenberger GA., Maler P.E., Tobler H.J., Wilson K., Monnier M.;
"The delta EEG (sleep)-inducing peptide (DSIP). XI. Amino-acid
                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
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Mammalia; Eutheria; Lagomorpha; Leporidae; Oryctolagus.
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Neoptera, Orthopteroidea, Dictyoptera, Blattaria, Blaberoidea,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 8;
Length 9;
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8 AA; 784 MW; 736365A5B9C865B8 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 16; DB 1; LA
Pred. No. 1.3e+05;
                            1.3e+05;
  DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (Rel. 01, Created)
(Rel. 01, Last sequence update)
(Rel. 40, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                 01-FEB-1991 (Rel. 17, Created)
01-FEB-1991 (Rel. 17, Last sequence update)
30-MAY-2000 (Rel. 39, Last annotation update)
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100.0%; Pred. No. 1...
0; Mismatches
  Score 17; DB 1
Pred. No. 1.3e+
0; Mismatches
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Leucophaea maderae (Madeira cockroach)
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P08939;
                                                                                                                             LCK2 LEUMA
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Best Local S
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                                                                                        LCK2_LEUMA
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TISSUE-Corpora cardiaca;

Machine A.K., Riley C.T., Fraser B.A., Nachman R.J.,

A Jaffe H., Raina A.K., Riley C.T., Fraser B.A., Nachman R.J.,

A Jaffe H., Raina A.K., Riley C.T., Fraser B.A., Nachman R.J.,

A Vogel V.W., Zhang Y.-S., Hayes D.K.;

Primary structure of two neuropeptide hormones with adipokinetic and

Thypotrehalosemic activity isolated from the corpora cardiaca of horse

THIS (Diptera).";

Proc. Natl. Acad. Sci. U.S.A. 86:8161-8164(1989).

THIS HORNOWE, RELEASED FROM CELLS IN THE CORPORA

CARDIACA AFTER THE BEGINNING OF FLIGHT, CAUSES RELEASE OF

DIGLYCERIDES FROM THE FAT BODY AND THEN STIMULATES THE FLIGHT

MUSCLES TO USE THESE DIGLYCERIDES AS AN ENERGY SOURCE.

SIMILARITY: BELONGS TO THE AKH / HRTH / RPCH FAMILY.
                                                                                                              MEDLINE=87175129; PubMed=3550726;
Graf M.V., Kastin A.J.;
Graf M.V., Kastin A.J.;
"Delta-sleep-inducing peptide (DSIP): an update.";
"Delta-sleep-inducing peptide (DSIP): an update.";
"Deptides 7:1165-1187(1296).
"IL PEPTIGES 7:1165-1187(1296).
"IL PEDLIGES 8:1187(1296).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Tabanus atratus (Horse fly).
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Brachycera; Tabanomorpha; Tabanidae;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
analysis, sequence, synthesis and activity of the nonapeptide."; Pflugers Arch. 376:119-129(1978).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             01-JAN-1990 (Rel. 13, Created)
01-FEB-1994 (Rel. 28, Last sequence update)
28-FEB-2003 (Rel. 41, Last annotation update)
Adipokinetic hormone (AKH) (Dipteran corpora cardiaca factor I)
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InterPro; IPR002047; AKH.
PROSITE; PS00256; AKH; 1.
Neuropeptide; Amidation; Flight; Pyrrolidone carboxylic acid.
PYRROLIDONE CARBOXYLIC ACID.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            h 33.3%; Score 16; DB 1; Length 9; Similarity 60.0%; Pred. No. 1.3e+05; 3; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          27.1%; Score 13; DB 1; Length 8; 100.0%; Pred. No. 1.3e+05; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0; Indels
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Best Local Similarity
Matches 2; Conserv
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Best Local Similarity
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Red pigment concentrating hormone (RPCH).

Pandalus borealis (Northern red shrimp).

Bukaryota; Metazoa; Arthropoda; Crustacea; Malacostraca;

Eumalacostraca; Bucarida; Decapoda; Pleocyemata; Caridea; Pandaloidea;

Pandalidae; Pandalus.
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Biochim. Biophys. Acta 371:304-311(1974).
-!- FUNCTION: THIS HORMONE ADAPTS THE ANIMAL TO LIGHT BACKGROUNDS BY
STIMULATING CONCENTRATION OF THE PIGMENT OF ITS RED BODY-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STIMULATES CONTRACTILE
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                                                                                                                                                                                                                                                                                                                                                                                                            neuropeptides
                         01-MAY-1991 (Rel. 18, Last sequence update)
01-MAY-1991 (Rel. 18, Last sequence update)
01-MAY-1991 (Rel. 18, Last annotation update)
Leucokinin II (L-II).
Leucophaea maderae (Madeira cockroach).
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Orthopteroidea; Dictyoptera; Blattaria; Blaberoidea;
NCBI TAXID=6988;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CHROMATOPHORES.
-!- SIMILARITY: BELONGS TO THE AKH / HRTH / RPCH FAMILY.
PIR; A61348; A61348.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   InterPro; IPR002047; AKH.
PROSITE; PS00256; AKH. 1.
Pigment; Hormone; Amidation; Pyrrolidone carboxylic
MOD_RES 1
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8 AA; 948 MW; 86786775B9C44736 CRC64;
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                                                                                                                                                                                                                                                                                                                                              TISSUB-Head;
Holman G.M., Cook B.J., Nachman R.J.;
"Isolation, primary structure and synthesis of tw
from Leucophaea maderae: members of a new family
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           27.1%; Score 13; DB 1; Le
100.0%; Pred. No. 1.3e+05;
ative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Comp. Biochem. Physiol. 84C.205-211(1986).
-!- FUNCTION: THIS CEPHALOMYOTROPIC PEPTIDE S'
ACTIVITY OF COCKROACH PROTODEUM (HINDGUT)
-!- SIMILARITY: TO THE OTHER LEUCOKININS.
NOU RES.
8 AMIDATION.
SEQÜENCE 8 AA; 852 MW; DC6365A5B9C8676A CJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               01-NOV-1988 (Rel. 09, Created)
01-FEB-1994 (Rel. 28, Last sequence update)
28-FEB-2003 (Rel. 41, Last annotation update)
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  8 AA
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STANDARD;
                                                                                                                                                                                                                                                                                                                          SEQUENCE, AND SYNTHESIS.
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Les 2; Conserv
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Eukaryota, Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha; Oestroidea;
Calliphoridae; Calliphora.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       wall components.
-!- DEVELOPMENTAL STAGE: Expressed throughout all walled developmental
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
  Gaps
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NCBI_TaxID=106964;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE, FUNCTION, SUBCELLULAR LOCATION, AND DEVELOPMENTAL STAGE MEDIINE=22044350; PubMed=12049410; Montes J.F., Durfort M., Ilado A., Garcia-Valero J.; Characterization and immunolocalization of a main proteinaceous component of the cell wall of the protozoan parasite Perkinsus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Calliphora vomitoriă.",
Proc. Natl. Acad. Sci. U.S.A. 89:2326-2330(1992).
-!- SIMILARITY: BELONGS TO THE FARP (FMRFAMIDE RELATED PEPTIDE)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TISSUE=Thoracic ganglion;
BUNDLINE=2191611; PubMed=1549595;
BUNC H., Johnsen A.H., Sewell J.C., Scott A.G., Orchard I.,
Rehfeld J.F., Thorpe A.;
Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 13; DB 1; Length 8;
Pred. No. 1.3e+05;
1; Mismatches 0; Indels
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(Rel. 32, Last sequence update)
(Rel. 32, Last annotation update)
                                                                                                                                                                                                                                                                                               Last sequence update)
Last annotation update)
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  Mismatches
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Best Local Similarity 66.7%;
Matches 2; Conservative
  Conservative
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                                                                                                                                                                                                                                                                      (Rel. 41,
                                                                                                                                                                                                                                                                                                                                               Wall protein-1 (PWP-1)
Perkinsus atlanticus.
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01-NOV-1995
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28-FEB-2003
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  5;
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-I- MISCELLANEOUS: ON THE 2D-GEL THE DETERMINED PI OF THIS UNKNOWN PROTEIN IS: 5.2, ITS MW IS: 50 kDa.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  "Plasma protein map: an update by microsequencing.";
Electrophoresis 13:707-714(1992).
-!- MISCELLANEOUS: ON THE 2D-GEL THE DETERMINED PI OF THIS UNKNOWN
PROTEIN IS: 7, ITS MW IS: 12 kDa.
                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Musinae; Mus.
NCBI_TaxID=10090;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Buteleostomi,
Mammalia, Butheria, Primates, Catarrhini, Hominidae, Homo.
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01-OCT-1994 (Rel. 30, Last sequence update)
01-FBE-1995 (Rel. 31, Last annotation update)
Unknown protein from 2D-page of fibroblasts (P50) (Fragment).
Mus musculus (Mouse)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   01-APR-1993 (Rel. 25, Created)
1-APR-1993 (Rel. 25, Last sequence update)
16-CT-2001 (Rel. 40, Last annotation update)
Unknown protein from 2D-page of plasma (Spot 36) (Fragment).
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Hughes G.J., Frutiger S., Paquet N., Ravier F., Pasquali
Sanchez J.-C., James R., Tissot J.-D., Bjellqvist B.,
Hochstrasser D.F.;
     Indels
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909 MW; 86677BS9D1A72042 CRC64;
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/FTId=VAR_000004.
                                                                                                                                                                                                                        8 A.
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  Mismatches
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  Conservative
                                                                                                                                                                                                                        STANDARD:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
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es 2; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
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P30096;
                                                                                                                                                                                                                      UF06 MOUSE
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SEQUENCE
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SEQUENCE
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SEQUENCE

P38644;

UFO6_MOUSE

RESULT 10

Query Match

Best Loc Matches

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UPAA_HUMAN

RESULT 11

SEQUENCE

VARIANT

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Search completed: November 17, 2003, 18:30:11
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Mustellus canis (Smooth dogfish).
Eukaryota; Metazoa; Chordaes; Craniata; Verrebrata; Chondrichthyes;
Elasmobranchii; Galeomorphii; Galeoidea; Carcharhiniformes; Triakidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Robey F.A., Tanaka T., Liu T.-Y.;
"Isolation and characterization of two major serum proteins from the dogfish, Mustelus canis, C-reactive protein and amyloid P component.";
J. Biol. Chem. 258:3889-3894(1983).
-!- SUBGUNIT: HOMOPENTAMEN. PENTAXIN (OR PENTRAXIN) HAVE A DISCOID ARRANGEMENT OF 5 NONCOVALENTLY BOUND SUBUNITS.
-!- SUBGEBLULAR LOCATION: Secreted.
-!- DISEASE: SAP IS A PRECURSOR OF AMYLOID COMPONENT P WHICH IS FOUND
                                                                                                                                                                                                                                                                                                                                                                                                                    TISSUE-Skin secretion;
Chen T.B., Orr D.F., Bjourson A.J., McClean S., Rao P.F., Shaw C.;
"Cloning and post-translational processing of frog skin kininogens.";
Submitted (ULL-2001) to the SMISS-PROT data bank.
-!- FUNCTION: [Thr6]bradykinin produces in vitro relaxation of rat
arterial smooth muscle and constriction of intestinal smooth
                                                                              Gaps
                                                                                                                                                                                                                                                                                                      [Thr6]bradykinin.
Bombina variegata (Yellow-bellied toad).
Elwaryota, Metachia, Chordata; Craniata; Vertebrata; Euteleostomi;
Amphibia; Batrachia; Anura; Archeobatrachia; Bombina.
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                                             27.1%; Score 13; DB 1; Length 9; 100.0%; Pred. No. 1.3e+05; Live 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE, SUBCELLULAR LOCATION, AND TISSUE SPECIFICITY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  -i- TISSUE SPECIFICITY: Skin.
-i- SIMILARITY: BELONGS TO THE BRADYKININ FAMILY.
Amphibian defense peptide; Vasodilator; Bradykinin.
SEQUENCE 9 AA; 1074 mW; 3339D771A9C86777 CRC64;
9 9 AMIDATION.
9 AA; 1068 MW; 39D10699CAB6D867 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-NOV-1990 (Rel. 16, Created)
01-NOV-1990 (Rel. 16, Last sequence update)
28-FEB-2003 (Rel. 41, Last annotation update)
Serum amyloid P-component (SAP) (Fragment).
                                                                                                                                                                                                                                                       28-FEB-2003 (Rel. 41, Created)
28-FEB-2003 (Rel. 41, Last sequence update)
15-SEP-2003 (Rel. 42, Last annotation update)
                                                                                                                                                                                                                           9 AA.
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NCBI_TaxID=7812;
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P83058;
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P19095;
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KNL3_BOMVA
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IN BASEMENT MEMBRANE AND ASSOCIATED WITH AMYLOID DEPOSITS.
-!-SIMILARITY: BELONGS TO THE PENTAXIN FAMILY.
PIR: B20569; B20569; B20569; PENTAXIN.
InterPro: IPR001759; Pentaxin.
PROSITE; PS00289; PENTAXIN; PARTIAL.
Lectin; Amyloid; Glycoprotein; Plasma; Pentaxin.

- 99
PENTAXIN.
                                                                                                                                                                                                                                                                                        0; Indels
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Best Local Similarity
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Title: Perfect score:

Sequence:

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Database

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Q99193 pseudomonas
Q99209 allus gall
Q92009 gallus gall
O12096 caprine art
O12100 caprine art
O12100 caprine art
O12104 caprine art
O1205 caprine art
O1205 caprine art
O12104 caprine art
O1206 caprine art
O1208 caprine art
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O64971 alfalfa mos
Q47410 escherichia
Q15999 homo sapien
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Q9gjv3 lagenorhync
Q9gjv1 lagenorhync
Q31415 gallus gall
Q98tuS xenopus lae
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TISSUE=TIBIAL GLAND;
Raftery M.J., Bradford A.M., Bowie J.H., Wallace J.C., Tyler M.J.;
"Peptides from Australian frogs. The structure of the dynastins from the banjo frogs Limnodynastes interioras, Limnodynastes dumerilii and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DYNASTIN 1.

Limnodynastes interioris (Giant banjo frog).

Limnodynastes interioris (Giant banjo frog).

Limnodynastes interioris (Giant banjo frog).

Limnodynastinas, Anura, Neobatrachia; Bufonoidea; Myobatrachidae;

Limnodynastinae; Limnodynastes.

NCBI_TAXID=30362;
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01-WAY-2000 (TrEMBLrel. 13, Last sequence update)
01-WAY-2000 (TrEMBLrel. 13, Last annotation update)
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80.0%; Pred. No. 8.3e+05;
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Aust. J. Chem. 46:833-842(1993).
-!- MASS SPECTROMETRY: MW=729; METHOD=FAB.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ALIGNMENTS
                                                                                                                                                                                                                                                       P93233
Q8GMM5
QQ9UMC7
P82685
Q9GM13
Q28866
Q60615
Q64971
Q47410
Q15999
Q9EQ02
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012096
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SEQUENCE 8 AA
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09twv0 anthopleura
QBaum7 carassius a
09hds4 aspergillus
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Q15901 homo sapien
O02032 lytechinus
Q9tt78 canis famil
Q9xsyl canis famil
Q95213 oryctolagus
Q81802 zea mays (m
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                    GenCore version 5.1.6
Copyright (c) 1993 - 2003 Compugen Ltd.
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Q9TWV0
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Q9S213
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Score

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Ikebe M., Hornick T., "Determination of the phosphorylation sites of smooth muscle caldesmon by protein kinase C."; Arch. Biochem. Biophys. 288:538-542(1991).
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Mammalia, Eutheria, Rodentia, Sciurognathi, Muridae, Murinae, Mus.
NCBI_TaxID=10090;
Oryctolagus cuniculus (Rabbit).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Lagomorpha; Leporidae; Oryctolagus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   "Stability of AML1 (core) site enhancer mutations in T-lymphomas induced by attenuated SL3-3 murine leukemia virus mutants."; J. Virol. 71:5080-5087(1997).
EMBL; Y11802; CAA72496.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
401-MAY-2000 (TrEMBLrel. 13, Last annotation update)
Anthor-RPAMIDE-NGUROPEPTIDE.
Anthor-PRAMIDE - Regantissima (Sea anemone).
Anthorpleura elegantissima (Sea anemone).
Bukaryota; Metazoa; Cnidaria; Anthoroa; Zoantharia; Actiniaria;
Nynamtheae; Actiniidae; Anthopleura.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Amtoft H.W., Sorensen A.B., Bareil C., Schmidt J., Luz A., Pedersen P.S.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 16; DB 11; Length 9;
Pred. No. 8.3e+05;
0; Mismatches 3; Indels
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SEQUENCE FROM N.A.
STRAIN=NMRI, TISSUE=Tumor;
MEDLINE=97332339; PubMed=9188573;
                                                             [1] =
SEQUENCE.
MEDLINE=91378498; PubMed=1898046;
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Similarity 57.1%;
4; Conservative (
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                                                                                                                                                                                                                                                             3; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mus musculus (Mouse).
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Best Local Similarity
Matches 4; Conserv
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Best Local Similarity
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                                             NCBI_TaxID=9986;
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01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-JUN-2002 (TrEMBLrel. 21, Last annotation update)
LOW density lipoprotein receptor-related protein (Fragment).
Gallus gallus (Chicken).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Archosauria; Aves; Neognathae; Galliformes; Phasianidae; Phasianinae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDLINE=92011685; PubMed=1918027;
Stifani S., Barber D.L., Aebersold R., Steyrer E., Shen X., Nimpf J.,
Schneider W.J.;
                                                                                                                                                                                                                                                                                                                                                                     Gaps
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                                                                                                                                                                                 Chang Y., Jeang K., Lietman T., Hayward G.S.; 
"Structural Organization of the Spliced Immediate-Early Gene Complex that Encodes the Major Acidic Nuclear (IE1) and Transactivator (IE2) 
Proteins of African Green Monkey Cytomegalovirus."; 
J. Biomed. Sci. 2:105-130(1995).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  "The laying hen expresses two different low density lipoprotein receptor related proteins.";
J. Biol. Chem. 266:19079-19087(1991).
NOW TER.
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                                                        Human herpesvirus 1.
Viruses; dsDNA viruses, no RNA stage; Herpesviridae;
Alphaherpesvirinae; Simplexvirus.
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01-%AXY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-UNN-2002 (TrEMBLrel. 21, Last annotation update)
CALDESMON=PHOSPHORYLATION site (Fragment).
                                                                                                                                                                                                                                                                                                         9 AA; 1029 MW; 797BB867740DDB04 CRC64;
                 01-MAR-2002 (TrEMBLrel. 20, Last annotation update)
[mmediate-early transactivator 110 (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               8 AA; 846 MW; C007272DD865BAAA CRC64;
   01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-MAR-2002 (TrEMBLrel. 20, Last annotation updat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         8 AA.
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Best Local Similarity 100.
Matches 3; Conservative
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Best Local Similarity
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                                                                                                                                                       STRAIN=MP;
PubMed=11725047;
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STRAIN=ST11;

MEDLINE=95047254; PubMed=7958782;

Constable A., Mollet B.;

"Isolation and characterisation of promoter regions from Streptococcus thermophilus.";

FEMS Microbiol. Lett. 122:85-90(1994).

FEMSL, X78210; CAA55045.1;

NON TER 8

SEQÜENCE 8 AA; 846 MW; ED086772D5B045B6 CRC64;
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                                                                                 3; Conservative
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01-DEC-2001 (TrEMBLrel.
STP6 protein (Fragment)
STP6.
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                                                                  Best Local Similarity
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NON TER 1
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 Polyprotein.
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            MEDLINE=93126143; PubMed=1480510; Carstensen K., Rinehart K.L., McFarlane I.D., Grimmelikhuijzen C.J.; Carstensen K., Kinehart K.L., McFarlane I.D., Grimmelikhuijzen C.J.; "Isolation of Leu-Pro-Pro-Gly-Pro-Leu-Pro-Arg-Pro-NH2 (Antho-RPamide), an N-terminally protected, biologically active neuropeptide from sea
                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE FROM N.A. Tishida M., Wood E., Callard G.V.; Tchoudakova A.V., Fishida M., Wood E., Callard G.V.; Promocter characteristics of two CYP19 genes differentially expressed in the brain and ovary of teleost fish.";
                                                                                                                                                  Gaps
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Actinopterygii; Neopterygii; Teleostei; Ostariophysi; Cypriniformes;
Cyprinidae; Carassius.
NCBI_TaxID=7957;
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Eukaryota; Fungi; Ascomycota; Pezizomycotina; Eurotiomycetes;
Eurotiales; Trichocomaceae; mitosporic Trichocomaceae; Aspergillus.
NCBI_TaxID=5059;
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Geiser D.M., Dorner J.W., Horn B.W., Taylor J.W.;
Geiser D.M., Dorner J.W., Horn B.W., Taylor J.W.;
The phylogenetics of mycotoxin and sclerotium production in
Aspergillus flavus and Aspergillus oryzae.";
Submitted (APR-2000) to the EMBL/GenBank/DDBJ databases.
EMBL; AF261861; AAG16135.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           31.2%; Score 15; DB 13; Length 9; 42.9%; Pred. No. 8.3e+05; Live 2; Mismatches 2; Indels
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                                                                                                                      Ouery Match 31.2%; Score 15; DB 5; Length 9; Best Local Similarity 75.0%; Pred. No. 8.3e+05; Matches 3; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                 01-WAR-2003 (TrEMBLrel. 23, Created)
01-WAR-2003 (TrEMBLrel. 23, Last sequence update)
01-WAR-2003 (TrEMBLrel. 23, Last annotation update)
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01-MAR-2001 (TrEMBLrel. 16, Last sequence update)
01-MAR-2001 (TrEMBLrel. 16, Last annotation update)
                                                                                    3:851-857(1992).
9 AA; 943 MW; 2908176737686777 CRC64;
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EMBL; AF324895; AAN32616.1; -.
EMBL; AF324896; AAN32617.1; -.
                                                                                                                                                                                                                                                                                                                                           Cytochrome P450 aromatase (Fragment)
                                                                                                                                                                                                                                                                        PRT;
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                                                                                                                                                                                                                                                                                                                                                                       Carassius auratus (Goldfish)
                                                                 anemones.";
Peptides 13:851-857(1992)
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Matches 3; Conservative
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09HDS4:
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"Alternative splicing of the sodium channel SCN8A predicts a truncated two-domain protein in fetal brain and non-neuronal cells.";
J. Biol. Chem. 272:24008-24015(1997).
EMBL, U97672; AAB80914.1;
MGD; MGI:103169; Scn8a.
                                                                                                                              Gaps
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Mammalia, Eutheria, Rodentia, Sciurognathi, Muridae, Murinae, Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-JAN-1998 (TrEMBLrel. 05, Created)
01-JAN-1998 (TrEMBLrel. 05, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
Truncated voltage-gated sodium channel alpha subunit (Fragment)
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                                                             29.2%; Score 14; DB 3; Length 8; 60.0%; Pred. No. 8.3e+05; tive 1; Mismatches 1; Indels
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Last annotation update)
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8 AA; 807 MW; F3B2C72AB5B87DD6 CRC64;
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SEQUENCE FROM N.A.
MEDLINE=97049323; PubMed=8894053;
Venta P.J., Brouillette J.A., Yuzbasiyan-Gurkan V., Brewer G.J.;
"Gene-specific universal mammalian sequence-tagged sites: application
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE FROM N.A. Vertander L., Yuzbasiyan-Gurkan V.; Venta P.J., Cao Y., Alexander L., Yuzbasiyan-Gurkan V.; "Dinucleotide repeat polymorphism in the canine retinoblastoma (RB1)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                MEDLINE=21015404; PubMed=11130975; Stroutllette J.A., Andrew J.R., Venta P.J.; Estimate of nucleotide diversity in dogs with a pool-and-sequence method.";
                                                                                                                                                                                                                                                                                                                         Canis familiaris (Dog).
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi;
Mammalia; Eutheria, Carnivora, Fissipedia, Canidae, Canis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Canis familiaris (Dog).
Sukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Carnivora, Fissipedia, Canidae, Canis.
NCBI_TaxID=9615,
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Submitted (JUN-1999) to the EMBL/GenBank/DDBJ databases.
EMBL; AF155737; AAD38807.1; -.
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01-NOV-1999 (TrEMBLrel. 12, Last sequence update)
01-NOV-1999 (TrEMBLrel. 12, Last annotation update)
Retinoblastoma protein (Fragment).
                                                                                                                                                                                                        01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
Thymidylate synthase (Fragment).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mamm, Genome 11:1079-1086(2000).
EMBL; AF202073; AAF20918.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    to the canine genome.";
Biochem. Genet. 34:321-341(1996).
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SEQUENCE FROM N.A.
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Q9XSY1
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TISSUBE-placenta,
TLEC C.-C., Yazdani A., Wehnert M., Bailey J., Couch L., Xiong M.,
Coolbaugh M.I., Chinault C.A., Baldini A., Lindsay E.A., Zhao Z.-Y.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Echinodermata; Eleutherozoa; Echinozoa;
Echinoidea; Euechinoidea; Echinacea; Temnopleuroida; Toxopneustidae;
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Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Caskey C.T.H.;
"Isolation of chromosome-specific genes by reciprocal probing of arrayed cDNAs and cosmid libraries.";
Hum. Mol. Genet. 0:0-0(1995).
EMBL.;__132080; AAA73891.1; -.
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Cserjesi P., Fang H., Brandhorst B.P.;
"Metallothionein gene expression in embryos of the sea urchin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 27.1%; Score 13; DB 5; Length 8; Best Local Similarity 100.0%; Pred. No. 8.38+05; Matches ?; Conservative 0; Mismatches 0; Indels
                                             Indels
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Length 8;
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01-JUL-1997 (TrEMBLrel. 04, Last sequence update)
01-NOV-1998 (TrEMBLrel. 08, Last annotation update)
Metallothionein (Fragment).
                                                                                                                                                                                                                                                                                                 01-NOV-1996 (TrEMBLrel. 01, Created)
01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
(Clone XP7B11B) (Fragment).
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DB 2; Le
8.3e+05;
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Query Match 27.1%; Score 13; DB Best Local Similarity 100.0%; Pred. No. 8.3 Matches 2; Conservative 0; Mismatches
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EMBL; U83400; AABS8320.1; -.
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STRAIN=F-I/rgm;
Mage R.G., Chen H.-T., Alexander C.B., Chen F.F.;
"Rabbit DOS2 and DH Gene Rearrangements in Early B-cell Development.";
Mol. Immunol. 0:0-0(1996).
BMBL; UG2585; AAB18735.1; -.
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Oryctolagus cuniculus (Rabbit).
Oryctolagus cuniculus (Rabbit).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Lagomorpha; Leporidae; Oryctolagus.
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27.1%; Score 13; DB 7; Length 8; Best Local Similarity 100.0%; Pred. No. 8.3e+05; Matches 2; Conservative 0; Mismatches 0; Indels
  0; Indels
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01-FEB-1997 (TrEMBLrel. 02, Created)
01-FEB-1997 (TrEMBLrel. 02, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
Germline DH (Df) gene (Fragment)
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SEQUENCE 8 AA; 845 MW; 5CA861B5AB58677B CRC64;
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  0; Mismatches
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Luteinising hormon GnRH peptide. Pet Peptide (221) inhi Mutant CD loop of Target fusion pept Fusion protein rel Mutant FG loop of Optional sequence #2 Linker peptide #3. Linker peptide #3. Linker peptide #1. Minimal motif #6. Gly-ala polymer of Linker peptide #1. Minimal motif #6. Gly-ala polymer of Linker peptide #1. Gly-ala polymer of Linker peptide #1. Gly-Ser tag peptid Peptide #3. Linker peptide #1. Gly-Ser tag peptid Peptide Modified humanised
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                                                                                   AAR86849
AAW94891
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AAB08104
AAB90979
AAB59836
AAR57721
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AAA026595
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99AU-0008073
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Decrein genes, and ABP00947-ABP01363 represent the proteins they encode.

Mag2732894-ABZ74687 represent human secreted protein genomic fragments. The invention also encompasses antibodies specific for the secreted proteins. The encompasses antibodies specific for the secreted proteins. The vectors and host cells comprishing a nucleic acid of the invention. The secreted proteins are thought to be involved in biological activities associated with cellular signalling, cellular differentiation, cell migration, prohormone activation and neurotransmitter activity. The secreted proteins, nucleic acids encoding them, antibodies or antibody fragments specific for the secreted proteins, and modulators of protein secreted proteins are useful for diagnosing or treating cancers or other hyperproliferative disorders. Additionally, the secreted proteins and thair nucleic acids may also be used in the treatment of autoimmune considers, inflammatory disorders, diseases involving angiogenesis, AIDS (acquired immunodeficiency syndrome), hepatitis, anaemia, and to promote cound healing. Nucleic acids of the invention may be used for chromosome individuals from minute biological samples, as hybridisation probes, and secreted protein of the invention.
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12-SEP-2001; 2001US-0950082.
12-SEP-2001; 2001US-0950083.
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                                                                                                                           The present sequence is a peptide fragment of human luteinising hormone releasing hormone (also known as LHRH, GnRH and gonadotrophin releasing hormone), which has spacers inserted at the N-terminus. It was used to demonstrate the novel adjuvant of the invention, which has lower reactogenicity then previous compositions. Vaccination of humans and animals against LHRH can be used as a method of fertility control, as well as enabling the control and treatment of disorders of the reproductive organs, such as testicular, breast, prostate and ovarian cancers, prostate enlargement and endometriosis. The composition of the linvention contains an amionic macromolecule and a saponin component, the latter of which is an immunostimulant, and it can also be used with other immunogens including soluble protein and it can also be used with other to a carrier protein and whole viruses.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New human secreted proteins and nucleic acids, useful for detecting or treating cancer or other hyperproliferative disorders, autoimmune disorders, inflammatory disorders, HIV disease, hepatitis or anemia -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; secreted protein; cancer; tumour; hyperproliferative disorder; autoimmune disorder; inflammation; angiogenic diseases; AIDS; acquired immunodeficiency syndrome; hepatitis; anaemia; wound healing; drug screening; chromosome identification; chromosome mapping; cytostatic; gene therapy; antiinflammatory; immunomodulator; anti-HIV;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human gene 284-encoded secreted protein HRDER22, SEQ ID NO:711.
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              can elicit an immune response in an animal, comprises an ionic polysaccharide component and a saponin component that is an
                                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%; Score 48; DB 21; Length 9; 100.0%; Pred. No. 9.3e+05; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 13; Page 1469; 2474pp; English.
                                                                                            Disclosure; Page 51; 53pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABR01230 standard; peptide; 9 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2001US-0950082
                                 polysaccharide component a immunostimulating complex
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Best Local Similarity luv.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     cytostatic; gene thera
antianaemic; vulnerary
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 GSGSGLRPG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GSGSGLRPG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           N-PSDB; ABZ73564.
                                                                                                                                                                                                                                                                                                                                                                                                                         9 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                12-SEP-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABR01230;
                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 2
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Human; secreted protein; nootropic; neuroprotective; cytostatic; virucide; dermatological; immunosuppressive; antiinflammatory; anti-HIV; vulnerary; antibacterial; antiparkinsonian; antisickling; antianaemic; antiartaritic; cancer; antirheumatic; hepatotropic; cerebroprotective; antiinflammatory; antiallergic; antidabetic; antiuleer; anticonvulsant, antiinflammatory; actiallergic; antidabetic; antiuleer; anticonvulsant, cardiovascular disorder; neurological disease; nephrotropic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New human secreted proteins encoded by genes contained in cDNA clones (e.g. HGCAC19), useful for preventing, treating or diagnosing e.g.
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60.4%; Score 29; DB 24; Length 9; 62.5%; Pred. No. 9.3e+05;
                           1; Indels
                           2; Mismatches
                                                                                                                                                                                                                                                Human secreted protein SEQ ID NO 684.
                                                                                                                                                            ABP99740 standard; Protein; 9 AA.
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1 GLRPG 5
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                                                                                                                                                                                                                                                                                                                                                                                                              5 AA;
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Modified-site
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20-MAR-1992;
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02-APR-1993
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AAR28241
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                                                                                           The invention relates to novel human genes (ABZ66891-ABZ68209) and the encoded secreted proteins (ABP99470-ABP99872) useful for preventing, treating or ameliorating medical conditions e.g. by protein or gene therapy. The genes are isolated from a range of human tissues disclosed in the specification. The nucleic acids, proteins, antibodies and (ant)agonists are useful in the diagnosis, treatment and prevention of: (a) cancer, e.g. breast and ovarian cancer and other cancers of the adrenal gland, bone, bone marrow, breast, gastrointestinal tract, liver, lung or urogenital; (b) immune disorders e.g. Addison's disease, altergies, autoimmune haemolytic anaemia, autoimmune thyroiditis, diabetes mellitus, Crohn's disease, multiple sclerosis, rheumatoid carthritis and ulcerative colitis; (c) cardiovascular disorders such as myocardial ischaemias; (d) wound healing; (e) neurological diseases e.g. cerebral anexia and epilepsy, and (f) infectious diseases such as viral,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CCK; neuropeptide; endorphin; hormone; LHRH; contraception; analgesia; improved bioavailability.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New alpha-substd. polypeptides are e.g. selective NK2 receptor
ligands - for treating inflammation, pain, stroke, ulcers,
hypertension, heart failure, depression, cancer, asthma, psychosis,
arthritis, etc.
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  AIDS, multiple sclerosis, herpes virus, leukemia, tick-borne encephalitis or West Nile fever
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 29; DB 24;
Pred. No. 9.3e+05;
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                                                                                                                                                                                                                                                                                                                                                                                          bacterial, fungal and parasitic infections
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                                                           Claim 1; Page 1469; 2423pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         62.5%;
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(first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 AGAGLPPG 8
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Best Local Similarity
Matches 5; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2 SGSGLRPG
                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 9 AA;
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Modified-site
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02-APR-1993
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generically claimed mono-, di, tri-, tetra- and penta-peptides
which include a substituent on an alpha-C atom in the chain. Such
substitution may modify the bioavailability, stability or
c absorbability of the peptide and hence may improve the activity of
the peptide as a drug. Depending on the nature of the parent peptide
(hormone, endorphin, CCK, NKZ, chemitacito peptide, etc.), the
modified peptides are variously useful for treating obesity, anxiety,
gastrointestinal ulcers, pain, stroke, inflammation, addictive drug
withdrawal symptoms, hypertension, heart failure, cognition or memory
discussion, psychosis and arthritis, and as contraceptives.
(Updated on 25-MAR-2003 to correct PD field.)
(Updated on 25-MAR-2003 to correct PR field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   endorphin; hormone; LHRH; contraception; analgesia;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New alpha-substd. polypeptides are e.g. selective NK2 receptor ligands - for treating inflammation, pain, stroke, ulcers, hypertension, heart failure, depression, cancer, asthma, psychosis, arthritis, etc.
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                                                       The peptide is a specifically claimed example of a group of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 5;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             58.3%; Score 28; DB 13; L
100.0%; Pred. No. 9.3e+05;
ive 0; Mismatches 0;
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Claim 3; Page 40; 46pp; English.
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92US-0852086
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(first entry)
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nes 5; Conservative
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The peptide is a specifically claimed example of a group of generically claimed mono., di., tri., tetra- and penta-peptides which include a substituent on an alpha-C atom in the chain. Such absorbability of the poptide and hence may improve the activity of the peptide as a drug. Depending on the nature of the parent peptide (hormone, endorphin, CCK, NK2, chemotactic peptide, etc.), the modified peptides are variously useful for treating obesity, anxiety, gastrointestinal ulcers, pain, stroke, inflammation, addictive drug withdrawal symptoms. Mypertension, heart failure, cognition or memory disorders, spasticity, depression, diabetes, cancer, asthma, bladder (Updated on 25-MAR-2003 to correct PM field.)

(Updated on 25-MAR-2003 to correct PM field.)

(Updated on 25-MAR-2003 to correct PM field.)
                                                                                                                                                                                                                                                                58.3%; Score 28; DB 13; Length 5; 100.0%; Pred. No. 9.3e+05; ive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                 AAR53131 standard; peptide; 5 AA.
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(first entry)
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Best Local Similarity
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                                                                                                                                                                                                                                          5 AA;
                                                                                                                                                                                                                                                                                                                                      GLRPG
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14-DEC-1994
                                                                                                                                                                                                                                            Sequence
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AAR53131
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                  The peptide is a specifically claimed example of a group of generically claimed mono-, di-, tri-, tetra- and penta-peptides which include a substituent on an alpha-C atom in the chain. Such absorbability of the program and hence may improve the activity of the peptide and hence may improve the activity of the peptide as a drug. Depending on the nature of the parent peptide (hormone, endorphin, CCK, NK2, chemotactic peptide, etc.), the modified peptides are variously useful for treating obseity, anxiety, gastrointestinal ulcers, pain, stroke, inflammation, addictive drug withdrawal symptoms, hypertension, heart failure, cognition or memory disorders, spaticity, depression, diabetes, cancer, asthma, bladder (Updated on 25-MAR-2003 to correct PV field.)

(Updated on 25-MAR-2003 to correct PV field.)

(Updated on 25-MAR-2003 to correct PV field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CCK; neuropeptide; endorphin; hormone; LHRH; contraception; analgesia; improved bioavailability.
                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New alpha-substd. polypeptides are e.g. selective NK2 receptor ligands - for treating inflammation, pain, stroke, ulcers, hypertension, heart failure, depression, cancer, asthma, psychosis,
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                                                                                                                                                                                                                                                                              58.3%; Score 28; DB 13; Length 5; 100.0%; Pred. No. 9.3e+05; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Howson W;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Hughes J, Richardson RS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /note= "alpha-Me-Pro"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                               AAR28242 standard; peptide; 5 AA.
 Claim 3; Page 41; 46pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Alpha-substituted short peptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /note= "Gly-NH2'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 3, Page 41, 46pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               91US-0690755.
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(first entry)
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                                                                                                                                                                                                                                                                                           Local Similarity 100.
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                                                                                                                                                                                                                                                                                                                                   5 GLRPG 9
                                                                                                                                                                                                                                                       5 AA;
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20-MAR-1992;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAR28242;
                                                                                                                                                                                                                                                        Sequence
                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                           Matches
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Gaps

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Peptide analogue, peptoid, cholecystokinin, CCK; obesity, anxiety, gastrointestinal ulcers, pain, stroke, inflammation, hypertension; heart failure, cognition; memory enhancement, spasticity, depression; diabetes; cancers, asthma; bladder dysfunction, psychosis, arthritis.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Richardson RS,
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Cholecystokinin analogue peptide #15.
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The sequences given in AAR53117-38 and AAR54530-51 are peptide analogues of cholecystokinin (CCK) which can be used to treat obesity, anxiety, gastrointestinal ulcers, pain, stroke, inflammation, hypertension, heart failure, cognition, memory enhancement, spasticity, depression, diabetes, cancers, asthma, bladder dysfunction, psychosis, arthritis
                                                                                                                                                                                                                                                                                                                                                      Peptide analogue; peptoid; cholecystokinin; CCK; obesity; anxiety; gastrointestinal ulcers; pain; stroke; inflammation; hypertension; heart failure; cognition; memory enhancement; spasticity; depression; diabetes; cancers; asthma; bladder dysfunction; psychosis; arthritis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                   Gaps
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Length 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Richardson RS
58.3%; Score 28; DB 15; 1 100.0%; Pred. No. 9.3e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              and in the treatment of substance withdrawl. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /note= "Amidated C-terminal"
                                   Mismatches
                                                                                                                                                                                                                                                                                                                      Cholecystokinin analogue peptide #17.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Hugues J,
 58.3.
100.0%; Pre
                                                                                                                                                                                           AAR53133 standard; peptide; 5 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /label= MePro
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(first entry)
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                                   Conservative
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Best Local Similarity
           Best Local Similarity
Matches 5, Conserv
                                                                  GLRPG 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      5 AA;
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Modified-site
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14-DEC-1994
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 Query Match
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(Updated on 25-MAR-2003 to correct PN field.)
The sequences given in AAR53117-38 and AAR54530-51 are peptide analogues
              of cholecystokinin (CCK) which can be used to treat obesity, anxiety, gastrointestinal ulcers, pain, stroke, inflammation, hypertension, heart failure, cognition, memory enhancement, spasticity, depression, diabetes, cancers, asthma, bladder dysfunction, psychosis, arthritis and in the treatment of substance withdrawl.
(Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Peptide analogue; peptoid; cholecystokinin; CCK; obesity; anxiety; gastrointestrial ulcers; pain; stroke; inflammation; hypertension; heart failure; cognition; memory enhancement; spasticity; depression; diabetes; cancers; asthma; bladder dysfunction; psychosie; arthritis.
                                                                                                                                                                                                                Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          C-terminal"
                                                                                                                                                                         58.3%; Score 28; DB 100.0%; Pred. No. 9.3 ive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Cholecystokinin analogue peptide #16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Howson W, Hugues J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /note= "Amidated
                                                                                                                                                                                                                                                                                                                                                                            AAR53132 standard; peptide; 5 AA
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                                                                                                                                                                                                              5; Conservative
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Best Local Similarity
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                                                                                                                                                                                                                                                                             1 GLRPG S
                                                                                                                                         5 AA;
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08-OCT-1993;
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                                                                                                                                         Sequence
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AAR53132
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RESULT 10

AAB90982

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The peptides of the invention are intermediates for the synthesis of gondorelin (luteinising hormone releasing hormone) and its analogues (see e.g. US 4024248).

(Updated on 16-AUG-2002 to add missing OS field.)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Prodn. of gonadorelin peptide intermediates without racemisation
- from new protected tryptophan tri:peptide derivs.
                                                                                   Gonadorelin; hormone; luteinising hormone releasing hormone.
                                                                                                                                                                                           'note= "bonded to urethane-protecting gp."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          58.3%; Score 28; DB 6; Length 8; 100.0%; Pred. No. 9.3e+05;
                                                    Sequence of gonadorelin peptide intermediate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    100.0%; Pred. ...
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                                                                                                                                                         Location/Qualifiers
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/note= "amidated"
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                                                                                                                                                                                                                              /label= Gly-NH2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 4; Page 23; 28pp; German.
                                                                                                                                                                                                                                                                                                                                                                       84DE-3411224.
                                                                                                                                                                                                                                                                                                                                      85EP-0103106
(updated)
(first entry)
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(first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                            Radscheit K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Immunogenic LHRH(3-10)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1985-243923/40.
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Best Local Similarity
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                                                                                                                                                            Key
Modified-site
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Modified-site
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16-AUG-2002
16-OCT-1991
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11-FEB-1993
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                                                                                                                                                                                                                                                                 EP156280-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                            Uhmann R,
                                                                                                                          Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic
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AAR26733
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention describes a modified therapeutic peptide (1) comprising a therapeutically active amino acid region (III) and a reactive group (II) (e.g. succinimidy) and maleimido groups) attached to a less therapeutically active amino acid region (IV), which covalently bonds with amino/hydroxyl/thiol groups on blood components to form a peptidase stabilised therapeutic peptide composed of 3-50 amino acids. (I) are useful for modifying therapeutic peptides e.g. hormones, growth factors and neurotransmitters, to protect them from peptidase activity in vivo for the treatment of various disorders. Endogenous therapeutic peptides are not suitable as drug candidates as they require frequent administration due to rapid degradation by peptidases in the body. Modifying and attaching therapeutic peptides to albumin prevents or reduces the action of peptidases to increase length of activity (half life) and specificity as bonding to large molecules decreases. AMB90829 to AAB92441 represent peptides which can be used in the exemplification of the present invention.
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                                                                                                                        Luteinising hormone releasing hormone (LH-RH) related peptide SEQ ID:156.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Modifying and attaching therapeutic peptides to albumin prevents peptidase degradation, useful for increasing length of in vivo activity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Gaps
                                                                                                                                                         Protection; endogenous therapeutic peptide; peptidase; conjugation; blood component; modification; succinimidyl; maleimido group; amino; hydroxyl; thiol; hormone; growth factor; neurotransmitter.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Thibaudeau K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              58.3%; Score 28; DB 22; Length 7; 100.0%; Pred. No. 9.3e+05; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Holmes DL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure; Page 241; 733pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Milner PG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAP50692 standard; peptide; 8 AA.
                    AAB90982 standard; Peptide; 7 AA.
                                                                                                                                                                                                                                                                                                                                                                                       99US-0134406.
99US-0153406.
99US-0159783.
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                                                                                     (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               (CONJ-) CONJUCHEM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2001-112059/12.
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Best Local Similarity
Matches 5; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              7 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GLRPG 7
                                                                                                                                                                                                                                                                                 WO200069900-A2
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10-SEP-1999;
15-OCT-1999;
                                                                                                                                                                                                                                  Homo sapiens
                                                                                       22-JUN-2001
                                                                                                                                                                                                                                                                                                                    23-NOV-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Bridon DP,
                                                                                                                                                                                                                                                  Synthetic.
                                                   AAB90982;
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RESULT 11

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AAP50692

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Gaps

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The invention relates immunogenic composition for eliciting an immune response to luteinising hormone releasing hormone (LHRH). The composition comprises a LHRH-diphtheria toxoid (DT) conjugate adsorbed to an ionic polysaccharide. The LHRH-OT compositions can be used for eliciting an immune response to LHRH, for castrating an animal, for regulating oestrus cycling in a female animal or for inhibiting characteristics induced by the sexual maturation of an animal, e.g. aggression or sexual activity. They can also be used for achieving production gains in livestock, e.g. reduction or elimination of unwanted organoleptic characteristics from the meat of livestock. They can also be used for inhibiting the growth of cells which are regulated directly or indirectly by LHRH, e.g. malignant oncofoetal cells or indirectly by LHRH, e.g. malignant concofoetal cells or cells, manignant ovarian cells, malignant concofoetal cells or of an animal. They can also be used for inhibiting pregnancy, prostate calargement, endometriosis or inflammatory responses The LHRH than the carrier adjuvant compositions. The effective immune response against LHRH results in prevention of the release of the hormones LH and an animal and an animal release of the hormones LH and animal and an animal and animal and animal animal and animal animal and animal anim
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                    New immunogenic leutenising hormone releasing hormone compositions comprise LHRH conjugated to diphtheria toxoid and adsorbed to an ionic polysaccharide, used to inhibit reproductive function in animals
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ..
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             58.3%; Score 28; DB 20; Length 8; 100.0%; Pred. No. 9.3e+05; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         endometriosis; prostate enlargement
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAB15364 standard; peptide; 8 AA.
                                                                                                                                                                                     Examples; Page 30; 41pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human LHRH peptide SEQ ID NO: 3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          99WO-AU01167
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity 100.00
Est Local Similarity 5; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             17-JAN-2001 (first entry)
WPI; 1999-120511/10.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          derivatives of LHRH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     σ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           8 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               LHRH(3-10) is highly immunogenic but lacks the hormonal properties of natural LHRH. Conjugates of the peptide with an immunogenic carrier protein can be used as an anti-LHRH vaccine. (An alpha-globulin/LHRH conjugate can also be used as anti-LHRH vaccine). The vaccines are administered shortly before slaughter to suppress the action of androgenic and non-androgenic hormones in non-castrated male animals. This allows the advantages associated with the male character (greater weight gain, more efficient feed utilisation and leaber carcasses) to be retained practically up to the time of slaughter. The treatment does not induce any local reactions which could result in the meat being rejected on grounds of quality.

(Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LHRH; immune response; luteinising hormone releasing hormone; DT; diphtheria toxoid; castrating; oestrus cycling; aggression; breast; sexual activity; organoleptic; livestock; cell growth; malignant; prostate; ovarian; oncofoetal; hyperplastic; pregnancy;
                                                                                                                                                                                                                                                                                                                                            steroid with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   58.3%; Score 28; DB 13; Length 8; 100.0%; Pred. No. 9.3e+05; rive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                        Improving meat quality of intact male animals - by immuno:neutralisation, shortly before slaughter, of anti-LHRH, esp. induced by two-stage vaccination
                                                                                                                                                                                                           Roulet C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                endometriosis; inflammatory response
                                                                                                                                                                                                           Dufour R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAW94892 standard; peptide; 8 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 22; Page 17; 18pp; French
                        92EP-0400496.
                                                                          91FR-0002513
                                                                                                  91FR-0015289
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Matches 5; Conservative
                                                                                                                                                    (INMR ) RHONE MERIEUX SA
                                                                                                                                                                                                        Chouvet C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LHRH peptide fragment.
                                                                                                                                                                                                                                                               WPI; 1992-294301/36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 8 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           5 GLRPG 9
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                     26-FEB-1992;
                                                                                                  10-DEC-1991;
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                                                                                                                                                                                                           Bonneau MB,
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Sequence

Query Match

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AAW94892;

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                                                                                 The present sequence is a peptide fragment of human luteinising hormone releasing hormone (also known as LHRH, GnRH and gonadotrophin releasing hormone). It was used to demonstrate the novel adjuvant of the invention, which has lower reactogenicity then previous compositions. Vaccination of humans and animals against LHRH can be used as a method of fertility control, as well as enabling the control and treatment of disorders of the reproductive organs, such as testicular, breast, prostate and ovarian cancers, prostate enlargement and endometriosis. The composition of the invention contains an anionic macromolecule and a saponin component, the latter of which is an immunostimulant, and it can also be used with other immunogens including soluble protein antigens, peptide haptens conjugated to a carrier protein and whole viruses.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Peptides of the SQs in AAP60174 and AAP60175 are claimed, the last 8 AAs
                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence of luteinising hormone releasing hormone (LHRH) analogue.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Contraception; vaccine; cryptorchidism; prostate cancer therapy;
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can elicit an immune response in an animal, comprises an ionic polysaccharide component and a saponin component that is an immunostimulating complex -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New nona- and deca-peptide(s) and dimers - are LHRH analogues
                                                                                                                                                                                                                                                                                                                                 DB 21; Length 8; 9.3e+05;
                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             useful for preventing ovulation and or treatment of cryptorchidism and prostate cancer
                                                                                                                                                                                                                                                                                                                                58.3%; Score 28; DB
100.0%; Pred. No. 9.3
:ive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Location/Qualifiers
                                                              Disclosure; Page 50; 53pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAP60174 standard; peptide; 9 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim li, Page 19; 20pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /label= Gly-NH2
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86AU-0057178.
86ZA-0003292.
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(first entry)
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                                                                                                                                                                                                                                                                                                                                                                                              5 GLRPG 9
                                                                                                                                                                                                                                                                                                    8 AA;
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19-AUG-1991
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                                                                                                                                                                                                                                                                                                                                                                  Matches
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of which are the same and in the same order as the last 8 residues of LHRH. Except at very low pH AAP60175 dimerises quickly through the SH gps. of Cys to give a dimer which is also claimed. A mixture of AAP60174 and AAP60175, and a conjugate of a carrier protein and 0.2-1.0 mg/kg given twice at a 3-6 week interval. (Updated on 25-MAR-2003 to correct PA field.)
                                                                                                                                                                                                           Gaps
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                                                                                                                                                                                                       0; Indels
                                                                                                                                                                        Length 9;
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100.0%; Pred. No. 9.3e+05;
ive 0; Mismatches 0;
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5; Conservative
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Best Local Similarity
Matches 5; Conserv
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                                                      US-09-192-854-117
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Sequence 154, Appl
Sequence 67, Appl
Sequence 14, Appl
Sequence 155, Appl
Sequence 48, Appl
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Sequence 204, App
Sequence 204, App
Sequence 48, Appl
Sequence 24, Appl
Sequence 47, Appl
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Sequence 4, Appli
Sequence 63, Appl
Sequence 99, Appl
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                                                                          November 17, 2003, 18:31:25; Search time 22.5 Seconds (without alignments) 73.024 Million cell updates/sec
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                                                                                                                                                                                                                                                                                                                                              | Cgn2_6/ptodata/2/pubpaa/US07_PUBCOMB.pep:*
| Cgn2_6/ptodata/2/pubpaa/US07_NEW_PUB.pep:*
| Cgn2_6/ptodata/2/pubpaa/US06_NEW_PUB.pep:*
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| Cgn2_6/ptodata/2/pubpaa/US06_PUBCOMB.pep:*
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| Cgn2_6/ptodata/2/pubpaa/US108_PUBCOMB.pep:*
| Cgn2_6/ptodata/2/pubpaa/US108_PUBCOMB.pep:*
| Cgn2_6/ptodata/2/pubpaa/US10_PUBCOMB.pep:*
         GenCore version 5.1.6
Copyright (c) 1993 - 2003 Compugen Ltd.
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US-09-968-561A-204

US-09-968-74A-204

US-09-968-74A-204

US-10-104-919-48

US-09-780-66BA-24

US-10-37-389-47

US-09-931-325A-154

US-09-931-325A-155

US-09-931-325A-155

US-09-947-249A-4

US-09-847-249A-4

US-09-847-249A-4

US-09-847-249A-4

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Maximum Match 100%
Listing first 45 summaries
                                                      - protein search, using sw model
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Gapop 10.0 , Gapext 0.5
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48
                                                                                                                                                1 GSGSGLRPG 9
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Match Length
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sequence 3, Appli
Sequence 117, Appli
Sequence 18, Appl
Sequence 63, Appl
               Sequence 17, Appl
Sequence 1000, Ap
Sequence 131, App
Sequence 278, App
Sequence 673, App
Sequence 131, App
Sequence 278, App
Sequence 278, App
Sequence 278, App
                                                                  Sequence 278, 1
Sequence 499, 1
Sequence 673, 2
Sequence 16, Ap
                                                                                                 Sequence 129, P
Sequence 45, P
Sequence 3, Ap
 Sequence 4
Sequence 7
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Sequence
Sequence
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US-10-082-747A-35
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ALIGNMENTS

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RESULT 2
US-09-968-561A-204
US-09-968-561A-204
Sequence 204, Application US/09968561A
Pacent No. US20020164642A1
GENERAL INFORMATION:
APPLICANT: TOMILISON, Ian M
APPLICANT: Winter, Gregory
TITLE OF INVENTION: Method to Screen Phage Display Libraries with Different Ligands
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
Sequence 117, Application US/09192854

Patent No. US20020068276A1

GENERAL INFORMATION

APPLICANT: Winter, Greg

APPLICANT: Tomlinson, Ian

TITLE OF INVENTION: Methods for Selecting Functional Peptides

FILE REFERENCE: 3789/72916

CURRENT APPLICATION NUMBER: US/09/192,854

CURRENT FILING DATE: 1998-11-17

EARLIER PEPLING DATE: 1997-11-21

NUMBER OF SEQ ID NOS: 2.22

SOFTWARE: PatentIn Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ..
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100.0%; Pre
0; 1
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Sequence 24 Application US/09780668A
; Sequence 24 Application US/09780668A
; Patent No. US20020147311A1
; GENERAL INPORMATION:
    APPLICANT: Gillies, Stephen
; APPLICANT: Lo, Kin-Wing
; TITLE OF INVENTION: Enhancing the Circulating Half-Life of Antibody-Based Fusion
; TITLE OF INVENTION: US/09/780,668A
; TITLE OF INVENTION: WINBER: US/09/780,668A
; FILE REFERENCE: LEX-01
; CURRENT FILING DATE: 2001-02-09
; PRIOR APPLICATION NUMBER: US 60/181,768
; SOFTWARE: PatentIn version 3.0
; SOFTWARE: PatentIn version 3.0
; TENDER OF SEQ ID NOS: 35
; SOS ID NO 24
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Sequence 47, Application US/10307389
Publication No. US20030175807A1
SEMERAL INFORMATION:
APPLICANT: BAUBET, VALERIE
APPLICANT: BRULET, PHEVE
APPLICANT: BRULET, PHILIPPE
TITLE OF INVENTION: CHIMERIC GFP-AEQUORIN AS BIOLUMINESCENT Ca++ REPORTERS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 15; Length 8; 6e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; OTHER INFORMATION: sequence at fusion junction US-09-780-668A-24
                                                                       APPLICANT: Chen, Zhi
APPLICANT: Chen, Zhi
APPLICANT: Hughes, Steven D.
TITLE OF INVENTION: Human Cytokine Receptor
FILE REFERENCE: 01-12
CURRENT APPLICATION NUMBER: US/10/104, 919
CURRENT FILING DATE: 2002-03-23
PRIOR FILING DATE: 2001-03-27
NUMBER: OF SEQ ID NOS: 62
SEQ ID NO 48
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 54.2%; Score 26; DB Best Local Similarity 100.0%; Pred. No. 6e+Matches 5; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; OTHER INFORMATION: Gly-Ser spacer peptide US-10-104-919-48
                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: PRT
ORGANISM: Artificial Sequence
                            Xu, Wenfeng
Kindsvogel, Wayne
Presnell, Scott R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 54.2'
Best Local Similarity 100.
Matches 5; Conservative
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US-09-780-668A-24
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TITLE OF INVENTION: Method to Screen Phage Display Libraries with Different Ligands
FILE REFERENCE: 8039/1073
CURRENT APPLICATION NUMBER: US/09/968,744A
CURRENT FILING DATE: 2003-01-13
PRIOR APPLICATION NUMBER: US 60/065,248
PRIOR PILING DATE: 1997-11-13
PRIOR FILING DATE: 1997-11-13
PRIOR FILING DATE: 1997-11-13
PRIOR FILING DATE: 1997-11-31
PRIOR FILING DATE: 1998-10-20
PRIOR FILING DATE: 1998-10-20
PRIOR FILING DATE: 200-02-24
NUMBER OF SEQ ID NOS: 350
SCOFTWARE: PATENTIN VENESION 3.1
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                 CURRENT PAPLICATION NUMBER: US/09/968,561A
CURRENT FILING DATE: 2001-10-01
PRIOR APPLICATION NUMBER: US 922131.1
PRIOR FILING DATE: 1997-10-20
PRIOR FILING DATE: 1997-10-20
PRIOR FILING DATE: 1997-11-13
PRIOR FILING DATE: 1997-11-21
PRIOR FILING DATE: 1997-11-21
PRIOR FILING DATE: 1997-11-21
PRIOR FILING DATE: 1998-10-20
PRIOR FILING DATE: 1998-10-20
PRIOR FILING DATE: 1998-10-20
PRIOR FILING DATE: 2000-02-4
NUMBER OF SEQ ID NOS: 350
SOFTWARE: PATENTING DATE: 2000-02-24
NUMBER OF SEQ ID NOS: 350
SOFTWARE: PATENTING DATE: 2000-02-24
NUMBER OF SEQ ID NOS: 350
SOFTWARE: PATENTING DATE: 2000-02-24
NUMBER OF SEQ ID NOS: 350
SOFTWARE: PATENTING DATE: 2000-02-24
NUMBER OF SEQ ID NOS: 350
SOFTWARE: PATENTING DATE: 2000-02-24
NUMBER OF SEQ ID NOS: 350
SOFTWARE: PATENTING DATE: 2000-02-24
NUMBER OF SEQ ID NOS: 350
SOFTWARE: PATENTING DATE: 2000-02-24
NUMBER OF SEQ ID NOS: 350
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US-09-968-744A-204
; Sequence 204, Application US/09968744A
; Publication No. US20030148372A1
; GENERAL INFORMATION:
; APPLICANT: Tomlinson, Ian M
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-10-344-919-48
; Sequence 48, Application US/10104919
; Publication No. US20030099608A1
; GENERAL INFORMATION:
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Matches 5; Conservative
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  FILE REFERENCE: 8039/1073B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
Matches 5; Conserv
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4 GLRPG 8
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RESULT 9
US-10-213-742-14
US-10-213-742-14

Sequence 14, Application US/10213742

Publication No. US20030194411A1

Publication No. US20030194411A1

GENERAL INFORMATION:

APPLICANT: Arye Rubinstein, Barry R. Bloom, Yair Devash and Stanley J. Cryz

TITLE OF INVENTION: PREVENTION OF HIV
                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/213,742
FILING DATE: 09-Aug-2002
CLASSIFICATION: <Unknown>
PRIOR APPLICATION NUMBER: US/08/946,525
FILING DATE: October 7, 1997
APPLICATION NUMBER: US/08/946,525
APPLICATION NUMBER: 08/785,696
                                TITLE OF INVENTION: Binding Phenol Oxidizing Enzyme-peptide TITLE OF INVENTION: Ending Phenol Oxidizing Enzyme-peptide FILE REFERENCE: GC690 CURRENT APPLICATION NUMBER: US/09/954,385 CURRENT FILING DATE: 2001-09-12 NUMBER OF SEQ ID NOS: 433 SOFFWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 67
                                                                                                                                                                                                                                                                                                                                                                        Score 24, DB 11; Length 7;
Pred. No. 6e+05;
0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE: Amster, Rothstein & Ebenstein
STREET: 90 Park Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEPHONE: (212) 697-5995
TELEFAX: (212) 286-0854 or 286-0082
TELEX: TWX 710-581-4766
INFORMATION FOR SEQ ID NO: 14:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE: January 17, 1997
ATTORNEY/AGENT INFORMATION:
NAME: Bogosian, Elizabeth A
REGISTRATION NUMBER: 39,911
REFERENCE/DOCKET NUMBER: 96700/448
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                          ; OTHER INFORMATION: binding peptide US-09-954-385-67
                       Winetzky, Deborah S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DESCRIPTION: peptide HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                             50.0%;
83.3%;
                                                                                                                                                                                                                                                                ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CITY: New York
STATE: New York
COUNTRY: U.S.A.
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Best Local Similarity 83..
Wang, Huaming
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     4 SGLRPG 9
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                                                                                                                                                                                                                                                                                                                                                                                                                       OTHER INFORMATION: Description of Artificial Sequence: Synthetic; OTHER INFORMATION: CONStruct US-10-307-389-47
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: MALARIA IMMUNGEN AND VACCINE
FILLE REFERENCE: 4564/83503 IGC-103.1
CURRENT APPLICATION NUMBER: US/09/931,325A
CURRENT FILING DATE: 2002-02-22
FRIOR APPLICATION NUMBER: 60/225,843
PRIOR APPLICATION NUMBER: 05.255,843
PRIOR FILING DATE: 2000-08-16
PRIOR FILING DATE: 2001-08-15
NUMBER OF SEQ ID NOS: 186
SOFTWARE: PATENTIN VET. 2.1
SEQ ID NO 154
AT THE SINGLE CELL LEVEL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3; Mismatches
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             FILE REFERENCE: 03495-0207-00000
CURRENT APPLICATION NUMBER: US/10/307,389
CURRENT FILING DATE: 2002-12-02
PRIOR APPLICATION NUMBER: US/09/863,901
PRIOR FILING DATE: 2001-05-24
PRIOR FILING DATE: 2000-06-01
PRIOR FILING DATE: 2000-06-01
PRIOR FILING DATE: 2000-06-06
PRIOR FILING DATE: 2000-06-06
PRIOR PILING DATE: 2000-12-14
NUMBER OF SEQ ID NOS: 48
SOFTWARE: PATENTIN VET: 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              54.2%; Score 26; 100.0%; Pred. No.
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APPLICANT: Baldwin, Toby D.
APPLICANT: Van Gastel, Franciscus J.C.
APPLICANT: Jansen, Giselle G.
APPLICANT: Murray, Christopher J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-09-931-325A-154
Sequence 154, Application US/09931325A
Publication No. US20030054337A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 67, Application US/09954385
Publication No. US20030100467A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               52.1%;
50.0%;
                                                                                                                                                                                                                                                                                                                                                        TYPE: PRT ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity 100.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity 50.0
Matches 4; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        47.9%; Score 23; DB 15; Length 6; 80.0%; Pred. No. 6e+05;
                                                                                                                                                                                                                                                                                                                           NAME: Trecartin, Richard F.
REGISTRATION NUMBER: 31,801
REFERENCE/DOCKET NUMBER: A-55186-11/RFT/BTC
TELECOMUNICATION INFORMATION:
TELEPHONE: 415-781-1989
           PRIOR APPLICATION DATA:

APPLICATION NUMBER: US/09/444,791
FILING DATE: 22-NO. US20030083464A1-1999
APPLICATION NUMBER: US 08/482,085
FILING DATE: 07-JUN-1995
APPLICATION NUMBER: US 08/175,155
FILING DATE: 29-DEC-1993
APPLICATION NUMBER: US 08/053,049
FILING DATE: 22-APR-1993
APPLICATION NUMBER: US 08/053,049
FILING DATE: 29-OCT-1897
APPLICATION NUMBER: US 07/114,618
FILING DATE: 29-OCT-1997
APPLICATION NUMBER: US 07/114,618
FILING DATE: 09-OCT-1997
APPLICATION NUMBER: US 07/114,618
FILING DATE: 09-OCT-1997
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PUBLICATION NO. US20030032588A1

GENERAL INFORMATION:

APPLICANT: WARSHALL, WILLIAM S.

APPLICANT: STARK, KEVIN LEE

TITLE OF INVENTION: GLUCAGON ANTACONIST

FILE REFERENCE: A-693
CURRENT APPLICATION NUMBER: US/09/847,249A

CURRENT FILING DATE: 2001-05-02

PRIOR PRDIAGONIST SOURCES OF PRIOR PRDIAGONIST SPIOR PRIOR DATE: 2000-05-03

NUMBER OF SEQ ID NOS: 80

SOFTWARE: PATENTIN VERSION 3.1

SSOFTWARE: PATENTIN VERSION 3.1

SSOFTWARE: PATENTIN VERSION 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY: linear
MOLECULE TYPE: peptide
SEQUENCE DESCRIPTION: SEQ ID NO: 48:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FEATURE:
COTHER INFORMATION: Preferred linker
US-09-847-249A-4
                                                                                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: 415-398-3249
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STRANDEDNESS: single
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ORGANISM: Artificial Sequence
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Best Local Similarity 80.0
Matches 4; Conservative
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GNGSG
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US-09-847-249A-4
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ADDRESSEE: Flehr Hohbach Test Albritton & Herbert LLP
STREET: Four Embarcadero Center, Suite 3400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 24; DB 11; Length 9; Pred. No. 6e+05; 3; Mismatches 1; Indels
                                                                Score 24; DB 12; Length 8; Pred. No. 6e+05;
                                                                                                            2; Indels
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
COMPUTER: PYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION UNMBER: US/10/096,986
> FILING DATE: 12-Mar-2002
                                                                                                                                                                                                                                              RESULT 10
US-09-931-325A-155
; Sequence 155, Application US/09931325A
; Bublication No. US20030054337A1
; GENERAL INFORMATION:
; APPLICANT: Birket, Ashley J.
; TITLE OF INVENTION: MALARIA IMMUNOGEN AND VACCINE
; FILE REFERENCE: 4564/48350 1CC-103.1
; CURRENT APPLICATION NUMBER: US/09/931,325A
; CURRENT FILING DATE: 2002-02-22
; PRIOR FILING DATE: 2000-08-16
; PRIOR FILING DATE: 2000-08-16
; NUMBER OF SEQ ID NOS: 186
; SOFIWARE: PALENTIN Ver. 2.1
; SEQ ID NO 155
; LENGTH: 9
                                                                                                               1; Mismatches
; SEQUENCE DESCRIPTION: SEQ ID NO: 14:
US-10-213-742-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 48, Application US/10096986; Publication No. US20030083464A1; GENERAL INFORMATION:
APPLICANT: Richardson, Charles Chambers, James Causey, Stuart Pollock, Thomas J. Cappello, Joseph; Caissman, John W. Crissman, John W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                50.0%;
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STATE: California
                                                              Query Match 50.0%;
Best Local Similarity 57.1%;
Matches 4; Conservative
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Best Local Similarity 50.0
Matches 4; Conservative
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Patent No. US20020002269A1
GENERAL INFORMATION:
APPLICANT: Milbrandt, Jeffrey D.
APPLICANT: Baloh, Robert H.
TITLE OF INVENTION: Artemin, A No. US20020002269A1el Neurotrophic Factor FILE REFERENCE: 6029-7996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 OTHER INFORMATION: Description of Artificial Sequence: canonical linker; OTHER INFORMATION: sequence variant
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APPLICANT: KLUG, Aaron
APPLICANT: KLUG, Aaron
APPLICANT: KLUG, Aaron
APPLICANT: KLUG, Aaron
APPLICANT: MOONE, Michael
TITLE OF INVENTION: FLEXIBLE LINKERS CONNECTED NUCLEIC ACID BINDING
TITLE OF INVENTION: PLEXIBLE LINKERS CONNECTED NUCLEIC ACID BINDING
TITLE OF INVENTION: MOLECULES
FILE REFERENCE: 8355-2011 / 411-031
CURRENT APPLICATION NUMBER: US/10/198,677
CURRENT FILING DATE: 2002-07-17
NUMBER OF SEQ ID NOS: 144
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Sequence 4, Application US/09840669B
Publication No. US20030040470A1
GENERAL INFORMATION:
I TILE OF INVENTION: APO-AI/AII PEPTIDE DERIVATIVES
FILE REFERENCE: A-69
CURRENT APPLICATION NUMBER: US/09/840,669B
CURRENT PILING DATE: 2002-06-07
PRIOR APPLICATION NUMBER: 60/198,920
PRIOR APPLICATION NUMBER: 60/198,920
PRIOR APPLICATION NUMBER: 60/198,920
PRIOR PILING DATE: 2000-04-21
NUMBER OF SEQ 1D NOS: 11
SOFTWARE: PatentIn version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                              CTHER INFORMATION: Preferred linker US-09-840-669B-4
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Publication No. US20030119023A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                           TYPE: PRT ORGANISM: Artificial Sequence
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Best Local Similarity 66.7
Matches 4; Conservative
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Best Local Similarity 80.0
Matches 4; Conservative
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2 GSGQKP 7
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2 GNGSG 6
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US-10-198-677-63
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LENGTH: 7
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LENGTH: 7
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MOLECULE TYPE: peptide
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OTHER INFORMATION:
OTHER INFORMATION:
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4, Appli
7, Appli
8, Appli
13, Appl
6, Appli
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81, Appl
18, Appl
18, Appl
19, Appl
52, Appl
52, Appl
3, Appli
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                                                                      November 17, 2003, 18:28:20 ; Search time 14.5 Seconds (without alignments) 26.262 Million cell updates/sec
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Sequence 13, 1
Sequence 6, Ap
Sequence 14, A
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Sequence 18,
Sequence 18,
Sequence 19,
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/cgn2_6/ptodata/1/iaa/5B_COMB.pep:*
/cgn2_6/ptodata/1/iaa/6A_COMB.pep:*
/cgn2_6/ptodata/1/iaa/6A_COMB.pep:*
/cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*
/cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*
GenCore version 5.1.6
(c) 1993 - 2003 Compugen Ltd.
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US-08-803-899-21
US-09-282-357-81
US-09-282-357-81
US-09-282-1908-18
US-08-29-1908-18
US-08-29-444-19
US-08-29-424-19
US-08-282-357-52
US-08-282-357-52
US-08-312-0718-12
US-08-51-656-17
US-08-312-0718-4
US-08-312-0718-4
US-08-312-0718-4
US-08-312-0718-4
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US-08-312-0718-8
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US-08-312-0718-8
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Maximum Match 100%
Listing first 45 summaries
                                                 OM protein - protein search, using sw model
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Gapop 10.0 , Gapext 0.5
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                                                                                                                          US-09-462-089-4
48
1 GSGSGLRPG 9
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Match Length
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                                                                                                                                         Perfect score:
                                                                                                                                                                               Scoring table:
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APPLICANT: Dufour, Raymond J.
APPLICANT: Dufour, Raymond J.
APPLICANT: Dufour, Claude J.M.
APPLICANT: Chouvet, Claire D.
APPLICANT: Bonneau, Michel B.
TITLE OF INVENTION: Method for improving the organoleptic
TITLE OF INVENTION: qualities of the meat from uncastrated male domestic
TITLE OF INVENTION: animals, vaccines which are usable in this method, new
TITLE OF INVENTION: peptide, in particular for producing these vaccines.
NUMBER OF SEQUENCES: 2
ADDRESSES: Larson and Taylor
     48, Appl
55, Appl
6, Appli
354, App
1000, Ap
99, Appl
25, Appl
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US-09-444-791A-48
US-08-529-190B-55
US-08-644-818-354
US-08-044-818-354
US-09-082-2799-1000
US-09-115-38-99
US-08-154-7128-25
US-08-154-7128-25
US-08-154-7128-25
US-08-154-7128-22
US-08-154-7128-22
US-08-154-716-3
US-08-151-767A-27
US-08-751-767A-27
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727 Twenty-Third Street, South
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ALIGNMENTS
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Patent No. 5573767
GENERAL INFORMATION:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
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3Y: linear
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: Virginia
RY: USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 1
US-08-343-883-2
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/label= NH2 /note= "amidated glycine"

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GENERAL INFORMATION:
APPLICANT: DONAHOE,
PATRICIANT: DONAHOE,
APPLICANT: WANG, TONGWEN
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR ENHANCING
TITLE OF INVENTION: CELLULAR RESPONSE TO TGF-BETA LIGANDS
NUMBER OF SEQUENCES: 26
CORRESPONDENCES: 26
CORRESPONDENCES: STERNE, GOLDSTEIN & FOX P.L.L.C
STREET: 1100 NEW YORK AVENUE, SUITE 600
CITY: WASHINGTON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 54.2%; Score 26; DB 1; Length 6; Best Local Similarity 100.0%; Pred. No. 2.5e+05; Matches 5; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTY:

ZIP: 20005
COMPUTER: PLORPY disk
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/803,899
FTIING DATE: 02/21/1997
COMPUTER READBLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPAGE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/264,002
FILING DATE: 22-UN-1994
CLASSIFICATION: WINDRER: P.38,347
REFERENCE/OCKET NUMBER: P.38,347
REFERENCE/DOCKET NUMBER: P38,347
REFERENCE/DOCKET NUMBER: P38,347
REFERENCE/DOCKET NUMBER: P38,347
REFERENCE/OCKET NUMBER: G19/455-5100
TELEPHONE: G19/455-5100
TELEPHONE: G19/455-5100
TELEPHONE: G19/455-5100
TELEPHONE: G19/455-5100
SEQUENCE CHARACTERISTICS:
LENGTH: 6 amino acids
TELENGTH: 6 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: STEFFE, ERIC K.
REGISTRATION NUMBER: 36,688
REFERENCE/DOCKET NUMBER: 0609.4240001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 21, Application US/08803899
Patent No. 5912224
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE: 02/22/1996
CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: STEFFE. FPT
                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY: linear
MOLECULE TYPE: peptide
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ) NAME/KEY: Peptide
; LOCATION: 1..6
US-08-264-002-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 GSGSG 5
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US-08-803-899-21
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Patent No. 5559019

GENERAL INFORMATION:

APPLICANT: GUI, JIAN-FANG

APPLICANT: FUI, XIANG-DONG

TITLE OF INVENTION: NOVEL PROTEIN SERINE KINASE, SRPKI

NUMBER OF SEQUENCES: 17

CORRESPONDENCE ADDRESS:

ADDRESSEE: SPENSLEY HORN JUBAS & LUBITZ

STREET: 1880 Century Park East, Fifth Floor

CITY: Los Angeles

STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                              58.3%; Score 28; DB 1; Length 8; 100.0%; Pred. No. 2.5e+05; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 9;
                                                                                                                                                                                                      Luteinizing hormone-relaesing hormone (Li activity of some synthetic polypeptides. Fragments shorter than decapeptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 2
5519002-1
; Patent No. 5519002
; Patent No. 5519002
; TITLE OF INVENTION: METHOD AND COMPOSITION FOR
; PREVENTING CONCEPTION
; NUMBER OF SEQUENCES: 2
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/26,180
; FILING DATE: 01-MAR-1993
; REPLICATION NUMBER: 697,127
; FILING DATE: 08-MAY-1991
; APPLICATION NUMBER: 250,557
; FILING DATE: 29-SEP-1988
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 6; Le
2.5e+05;
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  Schally, A. V. Arimura, A.
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Best Local Similarity 100.0
Matches 5; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         5; Conservative
                                                                                                                                                                                        Yanaihara, N
                                              Carra
Redding, T.
                                                                                                     Konig, W.
Wissman, H.
                                                                                                                                             Jaeger, G.
Sandow, J.
                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
Matches 5; Conserv
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5 GLRPG 9
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                                                                                                                                                                                                                                            TITLE: Fr
JOURNAL:
VOLUME: 4
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                                                                                                   AUTHORS:
AUTHORS:
AUTHORS:
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AUTHORS:
TITLE: LI
AUTHORS:
AUTHORS:
AUTHORS:
AUTHORS:
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                                                                                                                                                                                                                            Sequence 81, Application US/09282357
Patent No. 6242580
GENERAL INFORMATION:
GENERAL INFORMATION:
TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
CORRESPONDENCE:
ADDRESSE:
ADDRESSE:
ADDRESSE:
MORRISON & FOERSTER
STREET:
CORRESPONDENCE ADDRESS:
ADDRESSES:
ADDRESSE
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Pred. No. 2.5e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY: USA
ZIP: 20006-1888
COMPUTER READBLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/282,357
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     54.2%; Score 100.0%; Pred. No. 2.5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 31, Application US/09235230; Patent No. 6559287; GENERAL INFORMATION: APPLICANT: Bennett, Kelly, APPLICANT: Wolff, Edith A.; APPLICANT: Aruffo, Alejandro A.; APPLICANT: Greenfield, Brad W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 6 amino acids
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Best Local Similarity 100
Matches 5; Conservative
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US-09-282-357-81
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                                    GSGSG
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                                                                                                                                                                                                                                                                                                                                                                                                                    0; Indels
                                                                                                                                                                                                                                                                                                                                                       54.2%; Score 26; DB 2; Length 6; 100.0%; Pred. No. 2.5e+05; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STREET: 2000 Pennsylvania Avenue, NW, suite 5500 CITY: Washington STATE: DC COUNTRY: USA ZIP: 20006-1888
COMPUTRY: USA ZIP: DISKELTE COMPUTER READABLE FORM: MEDIUM TYPE: DISKETTE COMPUTER: IBM Compatible OPERATING SYSTEM: DOS SOFTWARE: FastSEQ for Windows Version 2.0 CURRENT APPLICATION NUMBER: US/08/918,288
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
APPLICANT: BOIME, Irving
APPLICANT: MOYLE, William R.
TITLE OF INVENTION: SINGLE-CHAIN FORMS OF THE
TITLE OF INVENTION: GLYCOPROTEIN HORMONE QUARTET
NUMBER OF SEQUENCES: 83
CORRESPONDENCE ADDRESS:
ADDRESSEE: MORRISON & POERSTER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: Murashige, Kate H
REGISTRATION NUMBER: 29,959
REFRENCE/DOCKET NUMBER: 29500-20050.25
TELECOMMUNICATION INFORMATION:
TELEFONE: 202-887-1500
TELEFRAX: 202-887-0763
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/282,357
FILING DATE:
APPLICATION NUMBER: 08/853,524
FILING DATE: 09-MAY-1997
APPLICATION NUMBER: 08/199,382
FILING DATE: 18-FEB-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 81, Application US/08918288 Patent No. 6238890
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202)371-2600
TELEFAX: (202)371-2540
INFORMATION FOR SEQ ID NO: 21:
SEQUENCE CHARACTERISTICS:
LENGTH: 6 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
LENGTH: 6 amino acids
                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity 100.
Matches 5; Conservative
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                                                                                                                                                                                                                                      TOPOLOGY: linear
MOLECULE TYPE: peptide
US-08-803-899-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: amino acid
STRANDEDNESS: sir
                                                                                                                                                                              TYPE: amino acid STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               linear
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Sequence 19, Application US/09029424A
| Reach No. 6030795
| GENERAL INFORMATION:
| APPLICANT: Saitch, Masao
| APPLICANT: Miyacono, Kohei
| APPLICANT: Miyacono, Kohei
| APPLICANT: Ichijo, Hidenori
| TITLE OF INVENTION: NUCLEIC ACID MOLECULE ENCODING TGF RECEPTOR HAVING MODIFIED
| TITLE OF INVENTION: GROWTH INHIBITION, AND ITS USE
| FILE REFERENCE: L0461/7027
| CURRENT APPLICATION NUMBER: US/09/029,424A
                                                                                                                                                                                                                                         APPLICANT: DONAGNE, PATRICIA K.
APPLICANT: WANG, TONGWEN
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR ENHANCING
TITLE OF INVENTION: CELLULAR RESPONSE TO TGF-BETA LIGANDS
MUMBER OF SEQUENCES: 26
CORRESPONDENCE ADDRESS:
ADDRESSEE: STERNE, KESSLER, GOLDSTEIN & FOX P.L.L.C
STREET: 1100 NEW YORK AVENUE, SUITE 600
CITY: WASHINGTON
STATE: D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION DATA:
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/803,899
FILING DATE: 02/21/1997
CLASSIFICATION TOMAR: 514
PRIOR APPLICATION NUMBER: 60/012,054
FILING APPLICATION NUMBER: 60/012,054
FILING APPLICATION NUMBER: 60/012,054
FILING APPLICATION NUMBER: 60/012,054
ATTORNEY/AGENT INFORMATION:
NAME: STEFFE, BRIC K.
REGISTRATION NUMBER: 36,688
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 54.2%; Score 26; DB 2; L
100.0%; Pred. No. 2.5e+05;
iive 0; Mismatches 0;
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TELECOMMUNICATION: INFORMATION:
TELEPHONE: (202)371-2600
TELEPAX: (202)371-2540
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ZIP: 20005
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                US-08-803-899-18
; Sequence 18, Application US/08803899
Patent No. 5912224
; GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY: linear MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   USA .
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Best Local Similarity
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                              GSGSG
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US-09-029-424-19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 18, Application US/08529190B
Patent No. 5833931
GENERAL INFORMATION:
APPLICANT: Masucci, Maria G.
TITLE OF INVENTION:
CLYCINE OF CONFERRING INVISIBILITY TO THE IMMUNE SYSTEM
                                                                                                                                                                                                                                                                                                                                                                         ..
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FILE REFERENCE: ON0153aSequences
CURRENT APPLICATION NUMBER: US/09/235,230
CURRENT FILING DATE: 1999-01-21
PRIOR APPLICATION NUMBER: 60/072,416
PRIOR APPLICATION NUMBER: 60/072,416
PRIOR FILING DATE: 1998-01-24
NUMBER OF SEQ ID NOS: 32
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 31
LENGTH: 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
SOFTWARE: Wordperfect 6.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/529,190B
FILING DATE: 15-SEP-1995
CLASSIFICATION DATA:
PRIOR APPLICATION DATA:
FILING DATE: 10-APP-1995
FILING DATE: 01-SEP-1995
ATTORNEY/AGENT INFORMATION:
NAME: Williams, Ph.D., Kathleen A
REFERENCE/DOCKET NUMBER: 34,380
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES: 76
CORRESSPONDENCE ADDRESS:
ADDRESSEE: Banner & Witcoff, Ltd.
STREET: One Financial Center
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELECOMMUNICATION INFORMATION: TELEPHONE: 617-345-9100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               E: Diskette
IBM Compatible
SYSTEM: DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: 617-345-9111
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity 100.
Matches 5; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       8 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ZIP: 02111
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                        ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                      1 GSGSG 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2 GSGSG 6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CITY: Boston
STATE: MA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-08-529-190B-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-08-529-190B-18
                                                                                                                                                                                                                                                                               US-09-235-230-31
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APPLICANT: ROEDERN, ERICH G.
APPLICANT: KESSLER, HORST
APPLICANT: KUSCHER, BERNHARD
APPLICANT: KURSCHER, BERNHARD
APPLICANT: KUERND, MICHAEL
APPLICANT: KLENNER, THOMAS
TITLE OF INVENTION: USE OF D-GLUCOPHRANURONIC ACIDS AND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 8;
                                                                                                                                                                                                                                                                                                     E: MORRISON & FOERSTER
2000 Pennsylvania Avenue, NW, suite 5500
                                                                                                                                                       GENERAL INFORMATION:
APPLICANT: BOIME, Irving
APPLICANT: MOYLE, William R.
APPLICANT: MOYLE, WILLIAM R.
TITLE OF INVENTION: GLYCOPROTEIN HORMONE QUARTET
NUMBER OF SEQUENCES: 83
CORRESPONDENCE ADDRESS:
ADDRESSEE: ANDRESSE:
SIRBET: 2000 Pennsylvania Avenue, NW, suite 550
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
54.2%; Score 26; DB 3; La
Best Local Similarity 100.0%; Pred. No. 2.5e+05;
Matches 5; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY: USA
ZIP: .2006-1888
COMPUTER READBLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FESTESC for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/282,357
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            29500-20050.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FLIANC TATE:
CLASSIFICATION: 536
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/918,288
FILING DATE: 25 AUG-1997
APPLICATION NUMBER: 08/853,524
FILING DATE: 09-MAY-1997
APPLICATION NUMBER: 08/199,382
FILING DATE: 18-FEB-1994
ATTORNEY/AGENT INFORMATION:
NAME: MUTABALIGE, Kate H
REGISTRATION NUMBER: 29,959
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; Sequence 12, Application US/08332071B ; Patent No. 5556836
                                                                                                                Sequence 52, Application US/09282357 Patent No. 6242580
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELECOMMUNICATION INFORMATION TELEPHONE: 202-887-1500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
LENGTH: 8 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEX:
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: 202-887-0763
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             amino acid
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1 GSGSG 5
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TOPOLOGY:
US-09-282-357-52
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US-08-332-071B-12
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                                                                                                                                                                                                   54.2%; Score 26; DB 3; Length 8; 100.0%; Pred. No. 2.5e+05; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 52, Application US/08918288
| Patent No. 6238890 |
| GENERAL INFORMATION: GENERAL INFORMATION: APPLICANT: BOIME, Irving APPLICANT: BOYLE, William R. TITLE OF INVENTION: SINGLE-CHAIN FORMS OF THE TITLE OF INVENTION: GLYCOPROTEIN HORMONE QUARTET NUMBER OF SEQUENCES: 83 |
| CORRESPONDENCE ADDRESS: ADDRESSE: MORRISON & FOERSTER ADDRESSE: MORRISON & FOERSTER STREET: 2000 Pennsylvania Avenue, NW, suite 5500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         54.2%; Score 26; DB 3; Le
100.0%; Pred. No. 2.5e+05;
tive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE:
APPLICATION NUMBER: 08/853,524
FILING DATE: 09-MAY-1997
APPLICATION NUMBER: 08/1997
FILING DATE: 18-FEB-1994
ATTORNEY/AGENT INFORMATION:
NAME: MUTASAIGG, KATCH
REGISTRATION NUMBER: 29,959
REFERENCE/DOCKET NUMBER: 29,500-20050.25
TELECOMMUNICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OPERATING SYSTEM: DOS
SOFTWARE: FatsLED for Windows Version 2.0
SUGRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/918,288
              EARLIER APPLICATION NUMBER: PCT/GB96/02179
EARLIER FILING DATE: 1996-09-04
NUMBER OF SEQ ID NOS: 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/282,357
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT FILING DATE: 1998-04-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEPHONE: 202-887-1500
TELEFAX: 202-887-0763
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity 100.0
Matches 5; Conservative
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
FRAGTH: 8 amino acids
                                                                                                                                                                                                 Query Match
Best Local Similarity 100.
Matches 5; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                  TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      OPERATING SYSTEM:
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                                                                                                                                                                                                                                                                                                                                                                                      RESULT 11
US-08-918-288-52
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-08-918-288-52
                                                                                                                                                             US-09-029-424-19
                                                                            SEQ ID NO 19
LENGTH: 8
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1 GSGSG 5

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RESULT 15
US-08-318-837-30
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   THEIR DERIVATIVES FOR INCORPORATION IN PHARMACOLOGICALLY ACTIVE PEPTIDES AND THEIR SALIS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 7;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STATE: D.C.
COUNTRY: U.S.A.
ZIP: 20005-5701
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/576,039
FILING DATE: 21-DEC-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ANTIGENS AND ANTIBODIES FOR THE DETECTION OF COLLAGEN 1
                                                                                                                                                                                                                                                                               ZIP: 20005

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            52.1%; Score 25; DB 1; I 83.3%; Pred. No. 2.5e+05; tive 0; Mismatches 1;
TITLE OF INVENTION: THEIR DERIVATIVES FOR IN TITLE OF INVENTION: ACTIVE PEPTIDES AND THEIR UNMBER OF SEQUENCES: 17
CARESPONDENCE ADDRESS:
ADDRESSEE: CUSHMAN DARBY & CUSHMAN, L.L.P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CARGAT A PEDLICALION DATA

CARGAT A PEDLICALION DATA

FILING DATE: 01-NOV-1994

CLASSIFICATION NUMBER: US, 071B

REGISTRATION NUMBER: 35, 843

REGISTRATION NUMBER: 35, 843

REFERENCE/DOCKET NUMBER: 326/216933

TELEPHONE: 202-861-3000

TELEPHONE: 202-861-3000

TELEPHONE: 202-861-3000

TELEPHONE: 70-861-3000

TELEPHONE: 202-861-3000

TELEPHONE: 202-861-3000
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Patent No. 5821065
GENERAL INFORMATION:
APPLICANT: NASER, Werner
APPLICANT: DR GER, Brigitte
APPLICANT: ESSIG, Ulrich
APPLICANT: HUBER, Erasmus
TITLE OF INVENTION: DETECTION OF COLL
NUMBER OF SEQUENCES: 3
CORRESPONDENCE ADDRESS:
                                                                                                                                         STREET: 1100 NEW YORK AVENUE, N.W. CUTY: WASHINGTON STATE: D.C. COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 52.1
Best Local Similarity 83.3
Matches 5; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY: linear
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Sequence 30, Application US/08318837
Sequence 30, Application US/08318837
Setent No. 5981277
GENERAL INFORMATION:
APPLICANT: RRANSEN, LUCIA, DEVOS, KATHLEEN; VAN DE VOORDE,
APPLICANT: ANDRE; VAN HEUVERSWYN, HUGO
TITLE OF INVENTION: NEW POLYPEPTIDES AND PEPTIDES, NUCLEIC ACID
TITLE OF INVENTION: IMMUNOLOGY
NUMBER OF SEQUENCES: 53
CORRESPONDENCE ADDRESS:
ANDRESPONDENCE ADDRESS:
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Pred. No. 2.5e+05;
0; Mismatches 2; Indels
PRIOR APPLICATION DATA:
APPLICATION NUMBER: DE P 44 46 232.8
FILING DATE: 23-DEC-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: DE 195 03 146.6
FILING DATE: 01-FEB-1995
ATTORNEY, AGENT INFORMATION:
NAME: Murray, ROBert B.
REGISTRATION NUMBER: 22,980
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFWARE: ASCII
CURRENT APPLICATION DATA:
PAPLICATION NUMBER: US/08/318,837
FILING DATE: 13-0CT-1994
CLASSIFICATION: 800
PRIOR APPLICATION NUMBER: PCT/EP 93/01022
APPLICATION NUMBER: PCT/EP 93/01022
APPLICATION NUMBER: PCT/EP 93/01022
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 92.401.231.3
ATTONNEY/AGENT INFORMATION:
NAME: CHARLES A. MUSERLIAN
REGISTRATION NUMBER: 19,683
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BIERMAN AND MUSERLIAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ZIP: 10016
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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TELEPHONE: (212) 661-8000
                                                                                                                                                                                                              TELECOMMUNICATION INFORMATION TELEPHONE: (202) 638-5000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             52.1%;
71.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE: BIERMAN AND M
STREET: 600 THIRD AVENUE
CITY: NEW YORK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: (212) 661-8002
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                        TELEPHONE: (202)638-500
TELEPAX: (202)638-4810
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 52.1
Best Local Similarity 71.4
Matches 5; Conservative
                                                                                                                                                                                                                                                                                                                                    LENGTH: 8 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY: linear
MOLECULE TYPE: peptide
US-08-576-039-3
                                                                                                                                                                                                                                                                                                                                                         amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STATE: NEW YORK
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i LENGTH: 8 amino acids

i TYPE: anino acid

i STRANDEDNESS: single

i TOPOLOCY: lines: single

i TOPOLOCY: lines: peptide

i RAGMENT TYPE: peptide

i ORGANISM: Mouse, human

i ORGANISM: Mouse, human

i ORGANISM: Mouse, human

i ORGANISM: Wouse, human

i ORGANISM: Wouse, human

i ORGANISM: Wouse, human

i ORGANISM: Wouse, human

i ORGANISM: Mouse, human
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